Human Secreted Proteins

This application is a continuation-in-part of, and claims benefit under 35 U.S.C. [1] § 119(e) based on copending U.S. Provisional Application No. 60/278,650 filed on March 27, 2001. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending U.S. Utility Application No. 09/833,245, filed on April 12, 2001, and PCT International Application Serial No. US01/11988, filed on April 12, 2001. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06043, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/167,061, filed on November 23,/1999, and U.S. Provisional Application No. 60/124,146, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06012, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/166,989, filed on November 23, 1999, and U.S. Provisional Application No. 60/124,093, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06058, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,654, filed on December 3, 1999, and U.S. Provisional Application No. 60/124,145, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06044, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,661, filed on/December 3, 1999, and U.S. Provisional Application No. 60/124,099, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06059, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,622, filed on December 3, 1999, and U.S. Provisional Application No. 60/124,096, filed on March 12, 1999. This application

is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06042, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,663, filed on December 3, 1999, and U.S. Provisional Application No. 60/2/24,143, filed on March 12, 1999. This application is also a continuation-in-part of and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06014, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,665, filed on Décember 3, 1999, and U.S. Provisional Application No. 60/138,598, filed on June 11, 1999, and U.S. Provisional Application No. 60/124,095, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06013, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,662, filed on December 3, 1999, and U.S. Provisional Application No. 60/138,626, filed on June 11,1999, and U.S. Provisional Application No. 60/125,360, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06049, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,667, filed on December 3, 1999, and U.S. Provisional Application No. 60/138,574, filed on June 11, 1999, and U.S. Provisional Application No. 60/124,144, filed on March 12, 1999. /This application is also a continuation-in-part of, and claims benefit under 35 U.S.C,/§ 120 of copending PCT International Application Serial No. US00/06057, filed on/March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,666, filed on December 3, 1999, and U.S. Provisional Application No. 60/138,597, filed on June 11, 1999, and U.S. Provisional Application No. 60/124,142, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06824, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,664, filed on December 3, 1999, and U.S. Provisional Application No. 60/125,359, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06765, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,906, filed on December 10, 1999, and

U.S. Provisional Application No. 60/126,051, filed on March 23, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06792, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,980, filed on December 10, 1999, and U.S. Provisional Application No. 60/125,362, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06830, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,910, filed on December 10, 1999, and U.S. Provisional Application No. 60/125,361, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. U\$00/06782, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,936, filed on December 10, 1999, and U.S. Provisional Application No. 60/125,812, filed on March 23, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06822, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,916, filed on December 10, 1999, and U.S. Provisional Application No. 60/126,054, filed on March/23, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06791, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) başed on U.S. Provisional Application No. 60/169,946, filed on December 10, 1999, and U/S. Provisional Application No. 60/125,815, filed on March 23, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06828, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,616, filed on December 8, 1999, and U.S. Provisional Application No. 60/125,358, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under/35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06823, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,623, filed on December 8, 1999, and U.S. Provisional Application No. 60/125,364, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending

 PCT International Application Serial No. US00/06781, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,617, filed on December 8, 1999, and U.S. Provisional Application No. 60/125,363, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07505, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,410, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,502, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U/S.C. § 120 of copending PCT International Application Serial No. US00/07440, filed/on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S./Provisional Application No. 60/172,409, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,503, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07506, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,412, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,505, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07507, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,408, fixed on December 17, 1999, and U.S. Provisional Application No. 60/126,594, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07535, filed on March 22, 2000, which claims benefit under 35 U.S.C./§ 119(e) based on U.S. Provisional Application No. 60/172,413, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,511, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07525, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U,S. Provisional Application No. 60/171,549, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,595, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07534, filed on March 22, 2000, which clairs benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No.

60/171,504, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,598, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07483, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/171,552, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,596, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07526, filed on March 22, . 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/171,550, filed on December 22, 1/999, and U.S. Provisional Application No. 60/126,600, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07527, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/171,551, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,501, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 3\$ U.S.C. § 120 of copending/PCT International Application Serial No. US00/07/661, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application/No. 60/174,847, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,504, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial/No. US00/07579, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,853, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,509, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § /120 of copending PCT International Application Serial No. US00/07723, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/242,710, filed on October 25, 2000, and U.S. Provisional Application No. 60/174,852, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,506, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07724, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,850, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,510, filed on

March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14929, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,851, filed on January 7, 2000, and U.S. Provisional Application No. 60/138,573, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07722, filed on March 23,/2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,871, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,508, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07578, filed on March 23, 2000, which claims be nefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,872, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,507, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07726, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U/S. Provisional Application No. 60/174,877, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,597, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07677, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,064, filed on January 14, 2000, and U.S. Provisional Application No. 60/154,373, filed on September 17, 1999, and U.S. Provisional Application No. 60/126,601, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07725, filed on March 23, 2000, which claims benefit under 35 U.S. C. § 119(e) based on U.S. Provisional Application No. 60/176,063, filed on January 14, 2000, and U.S. Provisional Application No. 60/126,602, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S/C. § 120 of copending PCT International Application Serial No. US00/09070/filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,052, filed on January 14, 2000, and U.S. Provisional Application No. 60/128,695, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/08982, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,069, filed on January 14, 2000, and U.S. Provisional Application No. 60/128,696, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/08983, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(¢) based on U.S. Provisional Application No. 60/176,068, filed on January 14, 2000, and U.S. Provisional Application No. 60/128,703, filed on April 9, 1999. This/application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09067, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,929, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,697, filed on April 9, 1999. This application is also a continuation/in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09066, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,926, and U.S. Provisional Application No. 60/128,698, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09068, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/177,050, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,699, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/08981, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 1/19(e) based on U.S. Provisional Application No. 60/177,166, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,701, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/08980, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,930, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,700, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09071, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,931, filed on/January 20, 2000, and U.S. Provisional Application No. 60/128,694, filed on

April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09069/filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/177,049, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,702, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of/copending PCT International Application Serial No. US00/15136, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,629, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14926, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,628, filed on Júne 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14963, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,631, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/15135, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,632, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14934, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,599, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14933, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,572, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/15137, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,625, filed/on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14\(\delta\)28, filed on June 1, 2000, which claims benefit under 35 U.S.C. \(\frac{1}{2}\)19(e) based on U.S. Provisional Application No. 60/138,633, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14973, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/,138,630, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14964, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,627, filed on June 11, 1999. application is also a continuation-in-part of, and claims benefit under 35/U.S.C. § 120 of copending PCT International Application Serial No. US00/26376, filed on September 26, 2000, which claims benefit under 35 U.S.C. § 119(e) based/on U.S. Provisional Application No. 60/155,808, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C./§ 120 of copending PCT International Application Serial No. US00/26371, filed on September 26, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/155,804, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26324, filed on September 26, 2000, which claims benefit under 35 § 119(e) based on U.S. Provisional Application No. 60/155,807, filed on U.S.C. September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. U\$00/26323, filed on September 26, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/155,805, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26337, filed on September 26, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/155,866, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US01/13318, filed on April 27, 2001, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/212,142, filed on June 16, 2000, and U.S. Provisional Application No. 60/201,194, filed on May 2, 2000. Each of the above referenced PCT applications were published in the English language. /Each of the above referenced priority applications are hereby incorporated by reference in their entireties.

Field of the Invention

The present invention relates to novel proteins. More specifically, isolated nucleic acid molecules are provided encoding novel polypeptides. Novel polypeptides and antibodies that bind to these polypeptides are provided. Also provided are vectors, host cells, and recombinant and synthetic methods for producing human polynucleotides and/or polypeptides, and antibodies. The invention further relates to diagnostic and therapeutic methods useful for diagnosing, treating, preventing and/or prognosing disorders related to these novel polypeptides. The invention further relates to screening methods for identifying agonists and antagonists of polynucleotides and polypeptides of the invention. The present invention further relates to methods and/or compositions for inhibiting or enhancing the production and function of the polypeptides of the present invention.

Background of the Invention

- [3] Unlike bacterium, which exist as a single compartment surrounded by a membrane, human cells and other eukaryotes are subdivided by membranes into many functionally distinct compartments. Each membrane-bounded compartment, or organelle, contains different proteins essential for the function of the organelle. The cell uses "sorting signals," which are amino acid motifs located within the protein, to target proteins to particular cellular organelles.
- One type of sorting signal, called a signal sequence, a signal peptide, or a leader sequence, directs a class of proteins to an organelle called the endoplasmic reticulum (ER). The ER separates the membrane-bounded proteins from all other types of proteins. Once localized to the ER, both groups of proteins can be further directed to another organelle called the Golgi apparatus. Here, the Golgi distributes the proteins to vesicles, including secretory vesicles, the cell membrane, lysosomes, and the other organelles.
- [5] Proteins targeted to the ER by a signal sequence can be released into the extracellular space as a secreted protein. For example, vesicles containing secreted proteins can fuse with the cell membrane and release their contents into the extracellular space a process called exocytosis. Exocytosis can occur constitutively or after receipt of a triggering signal. In the latter case, the proteins are stored in secretory vesicles (or

secretory granules) until exocytosis is triggered. Similarly, proteins residing on the cell membrane can also be secreted into the extracellular space by proteolytic cleavage of a "linker" holding the protein to the membrane.

Thus there exists a clear need for identifying and using novel secreted polynucleotides and polypeptides. Identification and sequencing of human genes is a major goal of modern scientific research. For example, by identifying genes and determining their sequences, scientists have been able to make large quantities of valuable human "gene products." These include human insulin, interferon, Factor VIII, tumor necrosis factor, human growth hormone, tissue plasminogen activator, and numerous other compounds. Additionally, knowledge of gene sequences can provide the key to treatment or cure of genetic diseases (such as muscular dystrophy and cystic fibrosis).

Summary of the Invention

The present invention relates to novel secreted proteins. More specifically, isolated nucleic acid molecules are provided encoding novel secreted polypeptides. Novel polypeptides and antibodies that bind to these polypeptides are provided. Also provided are vectors, host cells, and recombinant and synthetic methods for producing human polynucleotides and/or polypeptides, and antibodies. The invention further relates to diagnostic and therapeutic methods useful for diagnosing, treating, preventing and/or prognosing disorders related to these novel polypeptides. The invention further relates to screening methods for identifying agonists and antagonists of polynucleotides and polypeptides of the invention. The present invention further relates to methods and/or compositions for inhibiting or enhancing the production and function of the polypeptides of the present invention.

Detailed Description

Polynucleotides and Polypeptides

Description of Table 1A

[8] Table 1A summarizes information concerning certain polypnucleotides and polypeptides of the invention. The first column provides the gene number in the

application for each clone identifier. The second column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence disclosed in Table 1A. Third column, the cDNA Clones identified in the second column were deposited as indicated in the third column (i.e. by ATCC Deposit Number and deposit date). Some of the deposits contain multiple different clones corresponding to the same gene. In the fourth column, "Vector" refers to the type of vector contained in the corresponding cDNA Clone identified in the second column. In the fifth column, the nucleotide sequence identified as "NT SEQ ID NO:X" was assembled from partially homologous ("overlapping") sequences obtained from the corresponding cDNA clone identified in the second column and, in some cases, from additional related cDNA clones. The overlapping sequences were assembled into a single contiguous sequence of high redundancy (usually three to five overlapping sequences at each nucleotide position), resulting in a final sequence identified as SEQ ID NO:X. In the sixth column, "Total NT Seq." refers to the total number of nucleotides in the contig sequence identified as SEQ ID NO:X." The deposited clone may contain all or most of these sequences, reflected by the nucleotide position indicated as "5' NT of Clone Seq." (seventh column) and the "3' NT of Clone Seq." (eighth column) of SEQ ID NO:X. In the ninth column, the nucleotide position of SEQ ID NO:X of the putative start codon (methionine) is identified as "5' NT of Start Codon." Similarly, in column ten, the nucleotide position of SEQ ID NO:X of the predicted signal sequence is identified as "5" NT of First AA of Signal Pep." In the eleventh column, the translated amino acid sequence, beginning with the methionine, is identified as "AA SEQ ID NO:Y," although other reading frames can also be routinely translated using known molecular biology techniques. The polypeptides produced by these alternative open reading frames are specifically contemplated by the present invention.

In the twelfth and thirteenth columns of Table 1A, the first and last amino acid position of SEQ ID NO:Y of the predicted signal peptide is identified as "First AA of Sig Pep" and "Last AA of Sig Pep." In the fourteenth column, the predicted first amino acid position of SEQ ID NO:Y of the secreted portion is identified as "Predicted First AA of Secreted Portion". The amino acid position of SEQ ID NO:Y of the last amino acid encoded by the open reading frame is identified in the fifteenth column as "Last AA of ORF".

- [10] SEQ ID NO:X (where X may be any of the polynucleotide sequences disclosed in the sequence listing) and the translated SEQ ID NO:Y (where Y may be any of the polypeptide sequences disclosed in the sequence listing) are sufficiently accurate and otherwise suitable for a variety of uses well known in the art and described further below. For instance, SEQ ID NO:X is useful for designing nucleic acid hybridization probes that will detect nucleic acid sequences contained in SEQ ID NO:X or the cDNA contained in the deposited clone. These probes will also hybridize to nucleic acid molecules in biological samples, thereby enabling a variety of forensic and diagnostic methods of the invention. Similarly, polypeptides identified from SEQ ID NO:Y may be used, for example, to generate antibodies which bind specifically to proteins containing the polypeptides and the secreted proteins encoded by the cDNA clones identified in Table 1A and/or elsewhere herein
- [11] Nevertheless, DNA sequences generated by sequencing reactions can contain sequencing errors. The errors exist as misidentified nucleotides, or as insertions or deletions of nucleotides in the generated DNA sequence. The erroneously inserted or deleted nucleotides cause frame shifts in the reading frames of the predicted amino acid sequence. In these cases, the predicted amino acid sequence diverges from the actual amino acid sequence, even though the generated DNA sequence may be greater than 99.9% identical to the actual DNA sequence (for example, one base insertion or deletion in an open reading frame of over 1000 bases).
- [12] Accordingly, for those applications requiring precision in the nucleotide sequence or the amino acid sequence, the present invention provides not only the generated nucleotide sequence identified as SEQ ID NO:X, and the predicted translated amino acid sequence identified as SEQ ID NO:Y, but also a sample of plasmid DNA containing a human cDNA of the invention deposited with the ATCC, as set forth in Table 1A. The nucleotide sequence of each deposited plasmid can readily be determined by sequencing the deposited plasmid in accordance with known methods
- [13] The predicted amino acid sequence can then be verified from such deposits. Moreover, the amino acid sequence of the protein encoded by a particular plasmid can also be directly determined by peptide sequencing or by expressing the protein in a suitable host cell containing the deposited human cDNA, collecting the protein, and determining its sequence.

- [14] Also provided in Table 1A is the name of the vector which contains the cDNA plasmid. Each vector is routinely used in the art. The following additional information is provided for convenience.
- [15] Vectors Lambda Zap (U.S. Patent Nos. 5,128,256 and 5,286,636), Uni-Zap XR (U.S. Patent Nos. 5,128, 256 and 5,286,636), Zap Express (U.S. Patent Nos. 5,128,256 and 5,286,636), pBluescript (pBS) (Short, J. M. et al., *Nucleic Acids Res. 16:*7583-7600 (1988); Alting-Mees, M. A. and Short, J. M., *Nucleic Acids Res. 17:*9494 (1989)) and pBK (Alting-Mees, M. A. et al., *Strategies 5:*58-61 (1992)) are commercially available from Stratagene Cloning Systems, Inc., 11011 N. Torrey Pines Road, La Jolla, CA, 92037. pBS contains an ampicillin resistance gene and pBK contains a neomycin resistance gene. Phagemid pBS may be excised from the Lambda Zap and Uni-Zap XR vectors, and phagemid pBK may be excised from the Zap Express vector. Both phagemids may be transformed into *E. coli* strain XL-1 Blue, also available from Stratagene
- Vectors pSport1, pCMVSport 1.0, pCMVSport 2.0 and pCMVSport 3.0, were obtained from Life Technologies, Inc., P. O. Box 6009, Gaithersburg, MD 20897. All Sport vectors contain an ampicillin resistance gene and may be transformed into *E. coli* strain DH10B, also available from Life Technologies. See, for instance, Gruber, C. E., et al., *Focus* 15:59 (1993). Vector lafmid BA (Bento Soares, Columbia University, New York, NY) contains an ampicillin resistance gene and can be transformed into *E. coli* strain XL-1 Blue. Vector pCR[®]2.1, which is available from Invitrogen, 1600 Faraday Avenue, Carlsbad, CA 92008, contains an ampicillin resistance gene and may be transformed into *E. coli* strain DH10B, available from Life Technologies. See, for instance, Clark, J. M., *Nuc. Acids Res.* 16:9677-9686 (1988) and Mead, D. *et al.*, *Bio/Technology* 9: (1991).
- [17] The present invention also relates to the genes corresponding to SEQ ID NO:X, SEQ ID NO:Y, and/or a deposited cDNA (cDNA Clone ID). The corresponding gene can be isolated in accordance with known methods using the sequence information disclosed herein. Such methods include, but are not limited to, preparing probes or primers from the disclosed sequence and identifying or amplifying the corresponding gene from appropriate sources of genomic material.
- [18] Also provided in the present invention are allelic variants, orthologs, and/or species homologs. Procedures known in the art can be used to obtain full-length genes, allelic variants, splice variants, full-length coding portions, orthologs, and/or species homologs of genes corresponding to SEQ ID NO:X and SEQ ID NO:Y using information

from the sequences disclosed herein or the clones deposited with the ATCC. For example, allelic variants and/or species homologs may be isolated and identified by making suitable probes or primers from the sequences provided herein and screening a suitable nucleic acid source for allelic variants and/or the desired homologue.

[19] The present invention provides a polynucleotide comprising, or alternatively consisting of, the nucleic acid sequence of SEQ ID NO:X and/or a cDNA contained in ATCC Deposit No.Z. The present invention also provides a polypeptide comprising, or alternatively, consisting of, the polypeptide sequence of SEQ ID NO:Y, a polypeptide encoded by SEQ ID NO:X, and/or a polypeptide encoded by a cDNA contained in ATCC deposit No.Z. Polynucleotides encoding a polypeptide comprising, or alternatively consisting of the polypeptide sequence of SEQ ID NO:Y, a polypeptide encoded by SEQ ID NO:X and/or a polypeptide encoded by the cDNA contained in ATCC Deposit No.Z, are also encompassed by the invention. The present invention further encompasses a polynucleotide comprising, or alternatively consisting of the complement of the nucleic acid sequence of SEQ ID NO:X, and/or the complement of the coding strand of the cDNA contained in ATCC Deposit No.Z.

Description of Table 1B

[20] Table 1B summarizes some of the polynucleotides encompassed by the invention (including cDNA clones related to the sequences (Clone ID NO:Z), contig sequences (contig identifier (Contig ID:) and contig nucleotide sequence identifier (SEQ ID NO:X)) and further summarizes certain characteristics of these polynucleotides and the polypeptides encoded thereby. The first column provides the gene number in the application for each clone identifier. The second column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence disclosed in Table 1A and/or 1B. The third column provides a unique contig identifier, "Contig ID:" for each of the contig sequences disclosed in Table 1B. The fourth column provides the sequence identifier, "SEQ ID NO:X", for each of the contig sequences disclosed in Table 1A and/or 1B. The fifth column, "ORF (From-To)", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence of SEQ ID NO:X that delineate the preferred open reading frame (ORF) that encodes the amino acid sequence shown in the sequence listing and referenced in Table 1B as SEQ ID NO:Y (column 6). Column 7 lists residues comprising predicted epitopes contained in the polypeptides encoded by each of the preferred ORFs (SEQ ID NO:Y). Identification of potential

immunogenic regions was performed according to the method of Jameson and Wolf (CABIOS, 4; 181-186 (1988)); specifically, the Genetics Computer Group (GCG) implementation of this algorithm, embodied in the program PEPTIDESTRUCTURE (Wisconsin Package v10.0, Genetics Computer Group (GCG), Madison, Wisc.). This method returns a measure of the probability that a given residue is found on the surface of the protein. Regions where the antigenic index score is greater than 0.9 over at least 6 amino acids are indicated in Table 1B as "Predicted Epitopes". In particular embodiments, polypeptides of the invention comprise, or alternatively consist of, one, two, three, four, five or more of the predicted epitopes described in Table 1B. It will be appreciated that depending on the analytical criteria used to predict antigenic determinants, the exact address of the determinant may vary slightly. Column 8, "Tissue Distribution" shows the expression profile of tissue, cells, and/or cell line libraries which express the polynucleotides of the invention. The first number in column 8 (preceding the colon), represents the tissue/cell source identifier code corresponding to the key provided in Table 4. Expression of these polynucleotides was not observed in the other tissues and/or cell libraries tested. For those identifier codes in which the first two letters are not "AR", the second number in column 8 (following the colon), represents the number of times a sequence corresponding to the reference polynucleotide sequence (e.g., SEQ ID NO:X) was identified in the tissue/cell source. Those tissue/cell source identifier codes in which the first two letters are "AR" designate information generated using DNA array technology. Utilizing this technology, cDNAs were amplified by PCR and then transferred, in duplicate, onto the array. Gene expression was assayed through hybridization of first strand cDNA probes to the DNA array, cDNA probes were generated from total RNA extracted from a variety of different tissues and cell lines. Probe synthesis was performed in the presence of ³³P dCTP, using oligo(dT) to prime reverse transcription. After hybridization, high stringency washing conditions were employed to remove non-specific hybrids from the array. The remaining signal, emanating from each gene target, was measured using a Phosphorimager. Gene expression was reported as Phosphor Stimulating Luminescence (PSL) which reflects the level of phosphor signal generated from the probe hybridized to each of the gene targets represented on the array. A local background signal subtraction was performed before the total signal generated from each array was used to normalize gene expression between the different hybridizations. The value presented after "[array code]:" represents the mean of the duplicate values, following background subtraction and probe normalization. One of skill

in the art could routinely use this information to identify normal and/or diseased tissue(s) which show a predominant expression pattern of the corresponding polynucleotide of the invention or to identify polynucleotides which show predominant and/or specific tissue and/or cell expression. Column 9 provides the chromosomal location of polynucleotides corresponding to SEQ ID NO:X. Chromosomal location was determined by finding exact matches to EST and cDNA sequences contained in the NCBI (National Center for Biotechnology Information) UniGene database. Given a presumptive chromosomal location, disease locus association was determined by comparison with the Morbid Map, derived from Online Mendelian Inheritance in Man (Online Mendelian Inheritance in Man, OMIMTM. McKusick-Nathans Institute for Genetic Medicine, Johns Hopkins University (Baltimore, MD) and National Center for Biotechnology Information, National World Wide Web Medicine (Bethesda, MD) 2000. Library of http://www.ncbi.nlm.nih.gov/omim/). If the putative chromosomal location of the Query overlaps with the chromosomal location of a Morbid Map entry, an OMIM identification number is disclosed in column 10 labeled "OMIM Disease Reference(s)". A key to the OMIM reference identification numbers is provided in Table 5.

Description of Table 1C

Table 1C summarizes additional polynucleotides encompassed by the invention [21] (including cDNA clones related to the sequences (Clone ID NO:Z), contig sequences (contig identifier (Contig ID:) contig nucleotide sequence identifiers (SEQ ID NO:X)), and genomic sequences (SEO ID NO:B). The first column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence. The second column provides the sequence identifier, "SEQ ID NO:X", for each contig sequence. The third column provides a unique contig identifier, "Contig ID:" for each contig sequence. The fourth column, provides a BAC identifier "BAC ID NO:A" for the BAC clone referenced in the corresponding row of the table. The fifth column provides the nucleotide sequence identifier, "SEQ ID NO:B" for a fragment of the BAC clone identified in column four of the corresponding row of the table. The sixth column, "Exon From-To", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence of SEQ ID NO:B which delineate certain polynucleotides of the invention that are also exemplary members of polynucleotide sequences that encode polypeptides of the invention (e.g., polypeptides containing amino acid sequences encoded by the polynucleotide sequences delineated in column six, and fragments and variants thereof).

Description of Table 1D

- Table 1D: In preferred embodiments, the present invention encompasses a method of treating a disease or disorder listed in the "FEATURES OF PROTEIN" sections (below) and also as listed in the "Preferred Indications" column of Table 1D (below); comprising administering to a patient in which such treatment, prevention, or amelioration is desired a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) represented by Table 1A and Table 1D (in the same row as the disease or disorder to be treated is listed in the "Preferred Indications" column of Table 1D) in an amount effective to treat, prevent, or ameliorate the disease or disorder.
- [23] As indicated in Table 1D, the polynucleotides, polypeptides, agonists, or antagonists of the present invention (including antibodies) can be used in assays to test for one or more biological activities. If these polynucleotides and polypeptides do exhibit activity in a particular assay, it is likely that these molecules may be involved in the diseases associated with the biological activity. Thus, the polynucleotides or polypeptides, or agonists or antagonists thereof (including antibodies) could be used to treat the associated disease.
- The present invention encompasses methods of preventing, treating, diagnosing, or ameliorating a disease or disorder. In preferred embodiments, the present invention encompasses a method of treating a disease or disorder listed in the "Preferred Indications" column of Table 1D; comprising administering to a patient in which such treatment, prevention, or amelioration is desired a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) in an amount effective to treat, prevent, diagnose, or ameliorate the disease or disorder. The first and seccond columns of Table 1D show the "Gene No." and "cDNA Clone ID No.", respectively, indicating certain nucleic acids and proteins (or antibodies against the same) of the invention (including polynucleotide, polypeptide, and antibody fragments or variants thereof) that may be used in preventing, treating, diagnosing, or ameliorating the disease(s) or disorder(s) indicated in the corresponding row in Column 3 of Table 1D.
- [25] In another embodiment, the present invention also encompasses methods of preventing, treating, diagnosing, or ameliorating a disease or disorder listed in the "Preferred Indications" column of Table 1D; comprising administering to a patient

combinations of the proteins, nucleic acids, or antibodies of the invention (or fragments or variants thereof), sharing similar indications as shown in the corresponding rows in Column 3 of Table 1D.

- [26] The "Preferred Indication" column describes diseases, disorders, and/or conditions that may be treated, prevented, diagnosed, or ameliorated by a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof).
- [27] The recitation of "Cancer" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof) may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., leukemias, cancers, and/or as described below under "Hyperproliferative Disorders").
- In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D may be used for example, to diagnose, treat, prevent, and/or ameliorate a neoplasm located in a tissue selected from the group consisting of: colon, abdomen, bone, breast, digestive system, liver, pancreas, prostate, peritoneum, lung, blood (e.g., leukemia), endocrine glands (adrenal, parathyroid, pituitary, testicles, ovary, thymus, thyroid), uterus, eye, head and neck, nervous (central and peripheral), lymphatic system, pelvic, skin, soft tissue, spleen, thoracic, and urogenital.
- [29] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a pre-neoplastic condition, selected from the group consisting of: hyperplasia (e.g., endometrial hyperplasia and/or as described in the section entitled "Hyperproliferative Disorders"), metaplasia (e.g., connective tissue metaplasia, atypical metaplasia, and/or as described in the section entitled "Hyperproliferative Disorders"), and/or dysplasia (e.g., cervical dysplasia, and bronchopulmonary dysplasia).
- [30] In another specific embodiment, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a benign dysproliferative disorder selected from the group consisting of: benign tumors, fibrocystic conditions, tissue hypertrophy, and/or as described in the section entitled "Hyperproliferative Disorders".

- [31] The recitation of "Immune/Hematopoietic" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), blood disorders (e.g., as described below under "Immune Activity" "Cardiovascular Disorders" and/or "Blood-Related Disorders"), and infections (e.g., as described below under "Infectious Disease").
- In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having the "Immune/Hematopoietic" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: anemia, pancytopenia, leukopenia, thrombocytopenia, leukemias, Hodgkin's disease, non-Hodgkin's lymphoma, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, asthma, AIDS, autoimmune disease, rheumatoid arthritis, granulomatous disease, immune deficiency, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, immune reactions to transplanted organs and tissues, systemic lupus erythematosis, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, and allergies.
- [33] The recitation of "Reproductive" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the reproductive system (e.g., as described below under "Reproductive System Disorders").
- [34] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Reproductive" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cryptorchism, prostatitis, inguinal hernia, varicocele, leydig cell tumors, verrucous carcinoma, prostatitis, malacoplakia, Peyronie's disease, penile carcinoma, squamous cell hyperplasia, dysmenorrhea, ovarian adenocarcinoma, Turner's syndrome, mucopurulent cervicitis, Sertoli-leydig tumors, ovarian cancer, uterine cancer, pelvic inflammatory disease, testicular cancer, prostate cancer, Klinefelter's syndrome, Young's syndrome,

premature ejaculation, diabetes mellitus, cystic fibrosis, Kartagener's syndrome, testicular atrophy, testicular feminization, anorchia, ectopic testis, epididymitis, orchitis, gonorrhea, syphilis, testicular torsion, vasitis nodosa, germ cell tumors, stromal tumors, dysmenorrhea, retroverted uterus, endometriosis, fibroids, adenomyosis, anovulatory bleeding, amenorrhea, Cushing's syndrome, hydatidiform moles, Asherman's syndrome, premature menopause, precocious puberty, uterine polyps, dysfunctional uterine bleeding, cervicitis, chronic cervicitis, mucopurulent cervicitis, cervical dysplasia, cervical polyps, Nabothian cysts, cervical erosion, cervical incompetence, cervical neoplasms, pseudohermaphroditism, and premenstrual syndrome.

- [35] The recitation of "Musculoskeletal" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the immune system (e.g., as described below under "Immune Activity").
- [36] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Musculoskeletal" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: bone cancers (e.g., osteochondromas, benign chondromas, chondroblastoma, chondromyxoid fibromas, osteoid osteomas, giant cell tumors, multiple myeloma, osteosarcomas), Paget's Disease, rheumatoid arthritis, systemic lupus erythematosus, osteomyelitis, Lyme Disease, gout, bursitis, tendonitis, osteoporosis, osteoarthritis, muscular dystrophy, mitochondrial myopathy, cachexia, and multiple sclerosis.
- [37] The recitation of "Cardiovascular" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the cardiovascular system (e.g., as described below under "Cardiovascular Disorders").
- [38] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cardiovascular" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: myxomas,

fibromas, rhabdomyomas, cardiovascular abnormalities (e.g., congenital heart defects, cerebral arteriovenous malformations, septal defects), heart disease (e.g., heart failure, congestive heart disease, arrhythmia, tachycardia, fibrillation, pericardial Disease, endocarditis), cardiac arrest, heart valve disease (e.g., stenosis, regurgitation, prolapse), vascular disease (e.g., hypertension, coronary artery disease, angina, aneurysm, arteriosclerosis, peripheral vascular disease), hyponatremia, hypernatremia, hypokalemia, and hyperkalemia.

- [39] The recitation of "Mixed Fetal" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders").
- In specific embodiments, a protein, nucleic acid, or antibody of the invention [40] (or fragment or variant thereof) having a "Mixed Fetal" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: spina bifida, hydranencephaly, neurofibromatosis, fetal alcohol syndrome, diabetes mellitus, PKU, Down's syndrome, Patau syndrome, Edwards syndrome, Turner syndrome, Apert syndrome, Carpenter syndrome, Conradi syndrome, Crouzon syndrome, cutis laxa, Cornelia de Lange syndrome, Ellis-van Creveld syndrome, Holt-Oram syndrome, Kartagener syndrome, Meckel-Gruber syndrome, Noonan syndrome, Pallister-Hall syndrome, Rubinstein-Taybi syndrome, Scimitar syndrome, Smith-Lemli-Opitz syndrome, thromocytopenia-absent radius (TAR) syndrome, Treacher Collins syndrome, Williams syndrome, Hirschsprung's disease, Meckel's diverticulum, polycystic kidney disease, Turner's syndrome, and gonadal dysgenesis, Klippel-Feil syndrome, Ostogenesis imperfecta, muscular dystrophy, Tay-Sachs disease, Wilm's tumor, neuroblastoma, and retinoblastoma.
- [41] The recitation of "Excretory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and renal disorders (e.g., as described below under "Renal Disorders").

- [42] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Excretory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: bladder cancer, prostate cancer, benign prostatic hyperplasia, bladder disorders (e.g., urinary incontinence, urinary retention, urinary obstruction, urinary tract Infections, interstitial cystitis, prostatitis, neurogenic bladder, hematuria), renal disorders (e.g., hydronephrosis, proteinuria, renal failure, pyelonephritis, urolithiasis, reflux nephropathy, and unilateral obstructive uropathy).
- [43] The recitation of "Neural/Sensory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the nervous system (e.g., as described below under "Neural Activity and Neurological Diseases").
- [44] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Neural/Sensory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: brain cancer (e.g., brain stem glioma, brain tumors, central nervous system (Primary) lymphoma, central nervous system lymphoma, cerebellar astrocytoma, and cerebral astrocytoma, neurodegenerative disorders (e.g., Alzheimer's Disease, Creutzfeldt-Jakob Disease, Parkinson's Disease, and Idiopathic Presenile Dementia), encephalomyelitis, cerebral malaria, meningitis, metabolic brain diseases (e.g., phenylketonuria and pyruvate carboxylase deficiency), cerebellar ataxia, ataxia telangiectasia, and AIDS Dementia Complex, schizophrenia, attention deficit disorder, hyperactive attention deficit disorder, autism, and obsessive compulsive disorders.
- [45] The recitation of "Respiratory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the respiratory system (e.g., as described below under "Respiratory Disorders").

In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Respiratory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cancers of the respiratory system such as larynx cancer, pharynx cancer, trachea cancer, epiglottis cancer, lung cancer, squamous cell carcinomas, small cell (oat cell) carcinomas, large cell carcinomas, and adenocarcinomas. Allergic reactions, cystic fibrosis, sarcoidosis, histiocytosis X, infiltrative lung diseases (e.g., pulmonary fibrosis and lymphoid interstitial pneumonia), obstructive airway diseases (e.g., asthma, emphysema, chronic or acute bronchitis), occupational lung diseases (e.g., silicosis and asbestosis), pneumonia, and pleurisy.

The recitation of "Endocrine" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the respiratory system (e.g., as described below under "Respiratory Disorders"), renal disorders (e.g., as described below under "Renal Disorders"), and disorders of the endocrine system (e.g., as described below under "Endocrine Disorders".

In specific embodiments, a protein, nucleic acid, or antibody of the invention [48] (or fragment or variant thereof) having an "Endocrine" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cancers of endocrine tissues and organs (e.g., cancers of the hypothalamus, pituitary gland, thyroid gland, parathyroid glands, pancreas, adrenal glands, ovaries, and testes), diabetes (e.g., diabetes insipidus, type I and type II diabetes mellitus), obesity, disorders related to pituitary glands (e.g., hyperpituitarism, hypopituitarism, and pituitary dwarfism), hypothyroidism, hyperthyroidism, goiter, reproductive disorders (e.g. male and female infertility), disorders related to adrenal glands (e.g., Addison's Disease, corticosteroid deficiency, and Cushing's Syndrome), kidney cancer (e.g., hypernephroma, transitional cell cancer, and Wilm's tumor), diabetic nephropathy, interstitial nephritis, polycystic kidney disease, glomerulonephritis (e.g., IgM mesangial proliferative glomerulonephritis and glomerulonephritis caused by autoimmune disorders; such as Goodpasture's syndrome), and nephrocalcinosis.

- [49] The recitation of "Digestive" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the gastrointestinal system (e.g., as described below under "Gastrointestinal Disorders".
- In specific embodiments, a protein, nucleic acid, or antibody of the invention [50] (or fragment or variant thereof) having a "Digestive" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: ulcerative appendicitis, Crohn's disease, hepatitis, hepatic encephalopathy, portal hypertension, cholelithiasis, cancer of the digestive system (e.g., biliary tract cancer, stomach cancer, colon cancer, gastric cancer, pancreatic cancer, cancer of the bile duct, tumors of the colon (e.g., polyps or cancers), and cirrhosis), pancreatitis, ulcerative disease, pyloric stenosis, gastroenteritis, gastritis, gastric atropy, benign tumors of the duodenum, distension, irritable bowel syndrome, malabsorption, congenital disorders of the small intestine, bacterial and parasitic infection, megacolon, Hirschsprung's disease, aganglionic megacolon, acquired megacolon, colitis, anorectal disorders (e.g., anal fistulas, hemorrhoids), congenital disorders of the liver (e.g., Wilson's disease, hemochromatosis, cystic fibrosis, biliary atresia, and alpha1-antitrypsin deficiency), portal hypertension, cholelithiasis, and jaundice.
- The recitation of "Connective/Epithelial" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), cellular and genetic abnormalities (e.g., as described below under "Diseases at the Cellular Level "), angiogenesis (e.g., as described below under "Anti-Angiogenesis Activity "), and or to promote or inhibit regeneration (e.g., as described below under "Regeneration "), and wound healing (e.g., as described below under "Wound Healing and Epithelial Cell Proliferation").
- [52] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Connective/Epithelial" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of:

connective tissue metaplasia, mixed connective tissue disease, focal epithelial hyperplasia, epithelial metaplasia, mucoepithelial dysplasia, graft v. host disease, polymyositis, cystic disease, Alzheimer's dysplasia, tissue hypertrophy, hyperplasia, cerebral lymphoproliferative disorder, Waldenstron's macroglobulinemia, Crohn's disease, pernicious anemia, idiopathic Addison's disease, glomerulonephritis, bullous pemphigoid, Sjogren's syndrome, diabetes mellitus, cystic fibrosis, osteoblastoma, osteoclastoma, osteosarcoma, chondrosarcoma, osteoporosis, osteocarthritis, periodontal disease, wound nodosa, Wegener's polychondritis, vasculitis, polyarteritis relapsing healing, granulomatosis, cellulitis, rheumatoid arthritis, psoriatic arthritis, discoid lupus erythematosus, systemic lupus erythematosus, scleroderma, CREST syndrome, Sjogren's syndrome, polymyositis, dermatomyositis, mixed connective tissue disease, relapsing polychondritis, vasculitis, Henoch-Schonlein syndrome, erythema nodosum, polyarteritis nodosa, temporal (giant cell) arteritis, Takayasu's arteritis, Wegener's granulomatosis, Reiter's syndrome, Behcet's syndrome, ankylosing spondylitis, cellulitis, keloids, Ehler Danlos syndrome, Marfan syndrome, pseudoxantoma elasticum, osteogenese imperfecta, chondrodysplasias, epidermolysis bullosa, Alport syndrome, and cutis laxa.

Description of Table 2

Table 2 summarizes homology and features of some of the polypeptides of the [53] invention. The first column provides a unique clone identifier, "Clone ID NO:Z", corresponding to a cDNA clone disclosed in Table 1A or 1B. The second column provides the unique contig identifier, "Contig ID:" corresponding to contigs in Table 1B and allowing for correlation with the information in Table 1B. The third column provides the sequence identifier, "SEQ ID NO:X", for the contig polynucleotide sequence. The fourth column provides the analysis method by which the homology/identity disclosed in the Table was determined. Comparisons were made between polypeptides encoded by the polynucleotides of the invention and either a non-redundant protein database (herein referred to as "NR"), or a database of protein families (herein referred to as "PFAM") as further described below. The fifth column provides a description of the PFAM/NR hit having a significant match to a polypeptide of the invention. Column six provides the accession number of the PFAM/NR hit disclosed in the fifth column. Column seven, "Score/Percent Identity", provides a quality score or the percent identity, of the hit disclosed in columns five and six. Columns 8 and 9, "NT From" and "NT To" respectively, delineate the polynucleotides in "SEQ ID NO:X" that encode a polypeptide having a significant match to the PFAM/NR database as disclosed in the fifth and sixth columns. In specific embodiments polypeptides of the invention comprise, or alternatively consist of, an amino acid sequence encoded by a polynucleotide in SEQ ID NO:X as delineated in columns 8 and 9, or fragments or variants thereof.

Description of Table 3

Table 3 provides polynucleotide sequences that may be disclaimed according to [54] certain embodiments of the invention. The first column provides a unique clone identifier, "Clone ID", for a cDNA clone related to contig sequences disclosed in Table 1B. The second column provides the sequence identifier, "SEQ ID NO:X", for contig sequences disclosed in Table 1A and/or 1B. The third column provides the unique contig identifier, "Contig ID:", for contigs disclosed in Table 1B. The fourth column provides a unique integer 'a' where 'a' is any integer between 1 and the final nucleotide minus 15 of SEQ ID NO:X, and the fifth column provides a unique integer 'b' where 'b' is any integer between 15 and the final nucleotide of SEQ ID NO:X, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:X, and where b is greater than or equal to a + 14. For each of the polynucleotides shown as SEQ ID NO:X, the uniquely defined integers can be substituted into the general formula of a-b, and used to describe polynucleotides which may be preferably excluded from the invention. In certain embodiments, preferably excluded from the invention are at least one, two, three, four, five, ten, or more of the polynucleotide sequence(s) having the accession number(s) disclosed in the sixth column of this Table (including for example, published sequence in connection with a particular BAC clone). In further embodiments, preferably excluded from the invention are the specific polynucleotide sequence(s) contained in the clones corresponding to at least one, two, three, four, five, ten, or more of the available material having the accession numbers identified in the sixth column of this Table (including for example, the actual sequence contained in an identified BAC clone).

Description of Table 4

Table 4 provides a key to the tissue/cell source identifier code disclosed in Table 1B, column 8. Column 1 provides the tissue/cell source identifier code disclosed in Table 1B, Column 8. Columns 2-5 provide a description of the tissue or cell source. Codes corresponding to diseased tissues are indicated in column 6 with the word "disease". The use of the word "disease" in column 6 is non-limiting. The tissue or cell source may be

specific (e.g. a neoplasm), or may be disease-associated (e.g., a tissue sample from a normal portion of a diseased organ). Furthermore, tissues and/or cells lacking the "disease" designation may still be derived from sources directly or indirectly involved in a disease state or disorder, and therefore may have a further utility in that disease state or disorder. In numerous cases where the tissue/cell source is a library, column 7 identifies the vector used to generate the library.

Description of Table 5

Table 5 provides a key to the OMIM reference identification numbers disclosed [56] in Table 1B, column 10. OMIM reference identification numbers (Column 1) were derived from Online Mendelian Inheritance in Man (Online Mendelian Inheritance in Man, OMIM. McKusick-Nathans Institute for Genetic Medicine, Johns Hopkins University (Baltimore, MD) and National Center for Biotechnology Information, National 2000. World Wide Web Medicine, (Bethesda, MD) Library http://www.ncbi.nlm.nih.gov/omim/). Column 2 provides diseases associated with the cytologic band disclosed in Table 1B, column 9, as determined using the Morbid Map database.

Description of Table 6

[57] Table 6 summarizes some of the ATCC Deposits, Deposit dates, and ATCC designation numbers of deposits made with the ATCC in connection with the present application. These deposits were made in addition to those described in the Table 1A.

Description of Table 7

- [58] Table 7 shows the cDNA libraries sequenced, and ATCC designation numbers and vector information relating to these cDNA libraries.
- [59] The first column shows the first four letters indicating the Library from which each library clone was derived. The second column indicates the catalogued tissue description for the corresponding libraries. The third column indicates the vector containing the corresponding clones. The fourth column shows the ATCC deposit designation for each library clone as indicated by the deposit information in Table 6.

Definitions

- [60] The following definitions are provided to facilitate understanding of certain terms used throughout this specification.
- [61] In the present invention, "isolated" refers to material removed from its original environment (e.g., the natural environment if it is naturally occurring), and thus is altered "by the hand of man" from its natural state. For example, an isolated polynucleotide could be part of a vector or a composition of matter, or could be contained within a cell, and still be "isolated" because that vector, composition of matter, or particular cell is not the original environment of the polynucleotide. The term "isolated" does not refer to genomic or cDNA libraries, whole cell total or mRNA preparations, genomic DNA preparations (including those separated by electrophoresis and transferred onto blots), sheared whole cell genomic DNA preparations or other compositions where the art demonstrates no distinguishing features of the polynucleotide/sequences of the present invention.
- In the present invention, a "secreted" protein refers to those proteins capable of being directed to the ER, secretory vesicles, or the extracellular space as a result of a signal sequence, as well as those proteins released into the extracellular space without necessarily containing a signal sequence. If the secreted protein is released into the extracellular space, the secreted protein can undergo extracellular processing to produce a "mature" protein. Release into the extracellular space can occur by many mechanisms, including exocytosis and proteolytic cleavage.
- As used herein, a "polynucleotide" refers to a molecule having a nucleic acid sequence encoding SEQ ID NO:Y or a fragment or variant thereof (e.g., the polypeptide delinated in columns fourteen and fifteen of Table 1A); a nucleic acid sequence contained in SEQ ID NO:X (as described in column 5 of Table 1A and/or column 3 of Table 1B) or the complement thereof; a cDNA sequence contained in Clone ID NO:Z (as described in column 2 of Table 1A and/or 1B and contained within a library deposited with the ATCC); a nucleotide sequence encoding the polypeptide encoded by a nucleotide sequence in SEQ ID NO:B as defined in column 6 (EXON From-To) of Table 1C or a fragment or variant thereof; or a nucleotide coding sequence in SEQ ID NO:B as defined

in column 6 of Table 1C or the complement thereof. For example, the polynucleotide can contain the nucleotide sequence of the full length cDNA sequence, including the 5' and 3' untranslated sequences, the coding region, as well as fragments, epitopes, domains, and variants of the nucleic acid sequence. Moreover, as used herein, a "polypeptide" refers to a molecule having an amino acid sequence encoded by a polynucleotide of the invention as broadly defined (obviously excluding poly-Phenylalanine or poly-Lysine peptide sequences which result from translation of a polyA tail of a sequence corresponding to a cDNA).

In the present invention, "SEQ ID NO:X" was often generated by overlapping [64] sequences contained in multiple clones (contig analysis). A representative clone containing all or most of the sequence for SEQ ID NO:X is deposited at Human Genome Sciences, Inc. (HGS) in a catalogued and archived library. As shown, for example, in column 2 of Table 1B, each clone is identified by a cDNA Clone ID (identifier generally referred to herein as Clone ID NO:Z). Each Clone ID is unique to an individual clone and the Clone ID is all the information needed to retrieve a given clone from the HGS library. Table 7 provides a list of the deposited cDNA libraries. One can use the Clone ID NO:Z to determine the library source by reference to Tables 6 and 7. Table 7 lists the deposited cDNA libraries by name and links each library to an ATCC Deposit. Library names contain four characters, for example, "HTWE." The name of a cDNA clone (Clone ID) isolated from that library begins with the same four characters, for example "HTWEP07". As mentioned below, Table 1A and/or 1B correlates the Clone ID names with SEQ ID NO:X. Thus, starting with an SEQ ID NO:X, one can use Tables 1A, 1B, 6, 7, and 9 to determine the corresponding Clone ID, which library it came from and which ATCC deposit the library is contained in. Furthermore, it is possible to retrieve a given cDNA clone from the source library by techniques known in the art and described elsewhere herein. The ATCC is located at 10801 University Boulevard, Manassas, Virginia 20110-2209, USA. The ATCC deposits were made pursuant to the terms of the Budapest Treaty on the international recognition of the deposit of microorganisms for the purposes of patent procedure.

In specific embodiments, the polynucleotides of the invention are at least 15, at least 30, at least 50, at least 100, at least 125, at least 500, or at least 1000 continuous nucleotides but are less than or equal to 300 kb, 200 kb, 100 kb, 50 kb, 15 kb, 10 kb,

7.5kb, 5 kb, 2.5 kb, 2.0 kb, or 1 kb, in length. In a further embodiment, polynucleotides of the invention comprise a portion of the coding sequences, as disclosed herein, but do not comprise all or a portion of any intron. In another embodiment, the polynucleotides comprising coding sequences do not contain coding sequences of a genomic flanking gene (i.e., 5' or 3' to the gene of interest in the genome). In other embodiments, the polynucleotides of the invention do not contain the coding sequence of more than 1000, 500, 250, 100, 50, 25, 20, 15, 10, 5, 4, 3, 2, or 1 genomic flanking gene(s).

A "polynucleotide" of the present invention also includes those polynucleotides [66] capable of hybridizing, under stringent hybridization conditions, to sequences contained in SEQ ID NO:X, or the complement thereof (e.g., the complement of any one, two, three, four, or more of the polynucleotide fragments described herein), the polynucleotide sequence delineated in columns 7 and 8 of Table 1A or the complement thereof, the polynucleotide sequence delineated in columns 8 and 9 of Table 2 or the complement thereof, and/or cDNA sequences contained in Clone ID NO:Z (e.g., the complement of any one, two, three, four, or more of the polynucleotide fragments, or the cDNA clone within the pool of cDNA clones deposited with the ATCC, described herein), and/or the polynucleotide sequence delineated in column 6 of Table 1C or the complement thereof. "Stringent hybridization conditions" refers to an overnight incubation at 42 degree C in a solution comprising 50% formamide, 5x SSC (750 mM NaCl, 75 mM trisodium citrate), 50 mM sodium phosphate (pH 7.6), 5x Denhardt's solution, 10% dextran sulfate, and 20 μg/ml denatured, sheared salmon sperm DNA, followed by washing the filters in 0.1x SSC at about 65 degree C.

Also contemplated are nucleic acid molecules that hybridize to the polynucleotides of the present invention at lower stringency hybridization conditions. Changes in the stringency of hybridization and signal detection are primarily accomplished through the manipulation of formamide concentration (lower percentages of formamide result in lowered stringency); salt conditions, or temperature. For example, lower stringency conditions include an overnight incubation at 37 degree C in a solution comprising 6X SSPE (20X SSPE = 3M NaCl; 0.2M NaH₂PO₄; 0.02M EDTA, pH 7.4), 0.5% SDS, 30% formamide, 100 ug/ml salmon sperm blocking DNA; followed by washes at 50 degree C with 1XSSPE, 0.1% SDS. In addition, to achieve even lower stringency,

washes performed following stringent hybridization can be done at higher salt concentrations (e.g. 5X SSC).

- Note that variations in the above conditions may be accomplished through the inclusion and/or substitution of alternate blocking reagents used to suppress background in hybridization experiments. Typical blocking reagents include Denhardt's reagent, BLOTTO, heparin, denatured salmon sperm DNA, and commercially available proprietary formulations. The inclusion of specific blocking reagents may require modification of the hybridization conditions described above, due to problems with compatibility.
- [69] Of course, a polynucleotide which hybridizes only to polyA+ sequences (such as any 3' terminal polyA+ tract of a cDNA shown in the sequence listing), or to a complementary stretch of T (or U) residues, would not be included in the definition of "polynucleotide," since such a polynucleotide would hybridize to any nucleic acid molecule containing a poly (A) stretch or the complement thereof (e.g., practically any double-stranded cDNA clone generated using oligo dT as a primer).
- The polynucleotide of the present invention can be composed of any polyribonucleotide or polydeoxribonucleotide, which may be unmodified RNA or DNA or modified RNA or DNA. For example, polynucleotides can be composed of single- and double-stranded DNA, DNA that is a mixture of single- and double-stranded regions, single- and double-stranded RNA, and RNA that is mixture of single- and double-stranded regions, hybrid molecules comprising DNA and RNA that may be single-stranded or, more typically, double-stranded or a mixture of single- and double-stranded regions. In addition, the polynucleotide can be composed of triple-stranded regions comprising RNA or DNA or both RNA and DNA. A polynucleotide may also contain one or more modified bases or DNA or RNA backbones modified for stability or for other reasons. "Modified" bases include, for example, tritylated bases and unusual bases such as inosine. A variety of modifications can be made to DNA and RNA; thus, "polynucleotide" embraces chemically, enzymatically, or metabolically modified forms.
- [71] In specific embodiments, the polynucleotides of the invention are at least 15, at least 30, at least 50, at least 100, at least 125, at least 500, or at least 1000 continuous

nucleotides but are less than or equal to 300 kb, 200 kb, 100 kb, 50 kb, 15 kb, 10 kb, 7.5kb, 5 kb, 2.5 kb, 2.0 kb, or 1 kb, in length. In a further embodiment, polynucleotides of the invention comprise a portion of the coding sequences, as disclosed herein, but do not comprise all or a portion of any intron. In another embodiment, the polynucleotides comprising coding sequences do not contain coding sequences of a genomic flanking gene (i.e., 5' or 3' to the gene of interest in the genome). In other embodiments, the polynucleotides of the invention do not contain the coding sequence of more than 1000, 500, 250, 100, 50, 25, 20, 15, 10, 5, 4, 3, 2, or 1 genomic flanking gene(s).

"SEQ ID NO:X" refers to a polynucleotide sequence described in column 5 of Table 1A, while "SEQ ID NO:Y" refers to a polypeptide sequence described in column 10 of Table 1A. SEQ ID NO:X is identified by an integer specified in column 6 of Table 1A. The polypeptide sequence SEQ ID NO:Y is a translated open reading frame (ORF) encoded by polynucleotide SEQ ID NO:X. The polynucleotide sequences are shown in the sequence listing immediately followed by all of the polypeptide sequences. Thus, a polypeptide sequence corresponding to polynucleotide sequence SEQ ID NO:2 is the first polypeptide sequence shown in the sequence listing. The second polypeptide sequence corresponds to the polynucleotide sequence shown as SEQ ID NO:3, and so on.

The polypeptide of the present invention can be composed of amino acids [73] joined to each other by peptide bonds or modified peptide bonds, i.e., peptide isosteres, and may contain amino acids other than the 20 gene-encoded amino acids. polypeptides may be modified by either natural processes, such as posttranslational processing, or by chemical modification techniques which are well known in the art. Such modifications are well described in basic texts and in more detailed monographs, as well as in a voluminous research literature. Modifications can occur anywhere in a polypeptide, including the peptide backbone, the amino acid side-chains and the amino or carboxyl termini. It will be appreciated that the same type of modification may be present in the same or varying degrees at several sites in a given polypeptide. Also, a given polypeptide may contain many types of modifications. Polypeptides may be branched, for example, as a result of ubiquitination, and they may be cyclic, with or without branching. Cyclic, branched, and branched cyclic polypeptides may result from posttranslation natural processes or may be made by synthetic methods. Modifications include acetylation, acylation, ADP-ribosylation, amidation, covalent attachment of flavin, covalent attachment of a heme moiety, covalent attachment of a nucleotide or nucleotide derivative, covalent attachment of a lipid or lipid derivative, covalent attachment of phosphotidylinositol, cross-linking, cyclization, disulfide bond formation, demethylation, formation of covalent cross-links, formation of cysteine, formation of pyroglutamate, formylation, gamma-carboxylation, glycosylation, GPI anchor formation, hydroxylation, iodination, methylation, myristoylation, oxidation, pegylation, proteolytic processing, phosphorylation, prenylation, racemization, selenoylation, sulfation, transfer-RNA mediated addition of amino acids to proteins such as arginylation, and ubiquitination. (See, for instance, PROTEINS - STRUCTURE AND MOLECULAR PROPERTIES, 2nd Ed., T. E. Creighton, W. H. Freeman and Company, New York (1993); POSTTRANSLATIONAL COVALENT MODIFICATION OF PROTEINS, B. Johnson, Ed., Academic Press, New York, pgs. 1-12 (1983); Seifter et al., Meth. Enzymol. 182:626-646 (1990); Rattan et al., Ann. N.Y. Acad. Sci. 663:48-62 (1992)).

- "SEQ ID NO:X" refers to a polynucleotide sequence described, for example, in Tables 1A, 1B or 2, while "SEQ ID NO:Y" refers to a polypeptide sequence described in column 11 of Table 1A and or column 6 of Table 1B. SEQ ID NO:X is identified by an integer specified in column 4 of Table 1B. The polypeptide sequence SEQ ID NO:Y is a translated open reading frame (ORF) encoded by polynucleotide SEQ ID NO:X. "Clone ID NO:Z" refers to a cDNA clone described in column 2 of Table 1A and/or 1B.
- [75] "A polypeptide having functional activity" refers to a polypeptide capable of displaying one or more known functional activities associated with a full-length (complete) protein. Such functional activities include, but are not limited to, biological activity, antigenicity [ability to bind (or compete with a polypeptide for binding) to an anti-polypeptide antibody], immunogenicity (ability to generate antibody which binds to a specific polypeptide of the invention), ability to form multimers with polypeptides of the invention, and ability to bind to a receptor or ligand for a polypeptide.
- [76] The polypeptides of the invention can be assayed for functional activity (e.g. biological activity) using or routinely modifying assays known in the art, as well as assays described herein. Specifically, one of skill in the art may routinely assay secreted polypeptides (including fragments and variants) of the invention for activity using assays as described in the examples section below.

[77] "A polypeptide having biological activity" refers to a polypeptide exhibiting activity similar to, but not necessarily identical to, an activity of a polypeptide of the present invention, including mature forms, as measured in a particular biological assay, with or without dose dependency. In the case where dose dependency does exist, it need not be identical to that of the polypeptide, but rather substantially similar to the dose-dependence in a given activity as compared to the polypeptide of the present invention (i.e., the candidate polypeptide will exhibit greater activity or not more than about 25-fold less and, preferably, not more than about three-fold less activity relative to the polypeptide of the present invention).

TABLE 1A

Last AA of ORF	17	40	5	36	36	32	112	40	40
First Last AA of AA Secreted of Portion ORF		37		18	18	24	21	26	39
		36		17	17	23	70	25	38
First Last AA AA of of Sig Sig Pep Pep	1	-		-	-	-	-	-	-
AA SEQ D NO:	2608	2609	2610	2611	2612	2613	2614	2615	2616
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	27	309	386	197 2611	236	172	717	320	95
5' NT of Start Codon		309	386	197	236	172	717	320	95
3' NT of Clone Seq.	1256	1760	1529	2114	2158	443	1302	1174	916
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1	225	1	1	109	1	482	1	1
Total NT Seq.	1256	1760	1529	2114	2158	443	1315	1174	916
SEQ NO:	11	12	13	14	15	16	17	18	19
Vector	pBluescript SK-	pBluescript SK-	pBluescript SK-	pBluescript SK-	203917 pBluescript SK-04/08/99	203917 pBluescript SK-04/08/99	203917 pBluescript SK-04/08/99	pBluescript SK-	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917	203917 04/08/99		203917 04/08/99			203917 04/08/99	203917 04/08/99
cDNA Clone ID	H2CBD20	Н2СВН91	H2LBA54	H2LBB09	H2LBB09	H2MAC63	H2MBA76	H2MBF60	H6BSM88
Gene No.	_	2	8	4	5	9	7	∞	6

Last AA of ORF	246	27	111	47	48	25	104	19	83	104
First AA of Secreted Portion	21	5	35	28	25		22	8	20	36
	70	4	34	27	24		21	7	19	35
First Last AA AA of of Sig Sig Pep	-	-	1	-	1		-	1	1	1
AA SEQ ID NO: Y	2617	2618	2619	2620	2621	2622	2623	2624	2625	2626
5' NT AA of SEQ First ID AA of NO: Signal Y Pep	196	1434	190	175	98	275	329	683	1060	654
5' NT of of of Start AA of Codon Signal Pep	196		190	175	86	275	329		1060	654
3' NT of Clone Seq.	1228	1083	1397	951	229	508	1085	066	3152	1402
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	-		-	1	463	688	1
Total NT Seq.	1228	1960	1425	951	229	508	1099	066	3152	1402
SEQ NO:	20	21	22	23	24	25	26	27	28	29
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	PCRII	Other	pSport1	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203979 04/29/99	203918 04/08/99	PTA- 181 06/07/99	203917 04/08/99	203959	203917 04/08/99	PTA- 794 09/27/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	H6EEA48	H6EEN71	H6EEO05	H6EEU40	H7TDB54	H7TMB95	HAAAT06	HACAD42	HACBJ11	HACBS86
Gene No.	10	11	12	13	14	15	16	17	18	19

Last AA of ORF	59	36	59	51	27	15	3	32	39	126	35
First AA of Secreted Portion	8	26	26	31	14	15		29	25	32	32
	7	25	25	30	13	14		28	24	31	31
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1		1	1	_	1	1	1	1	1	1
AA SEQ ID NO:	2627	2628	2629	2630	2631	2632	2633	2634	2635	2636	2637
5	329	105	205	202	161	140	240	254	238	324	230
5' NT of Start Codon		105	205	202	161	140	240	254	238	324	230
	841	996	1005	464	839	1102	1112	2531	954	3342	620
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	-	1	1		-	1		-	1
	841	996	1005	464	839	1102	1112	2531	954	3342	620
NT SEQ ID NO:	30	31	32	33	34	35	36	37	38	39	40
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99	203979 04/29/99	203917 04/08/99
cDNA Clone ID	HACBT91	HACBZ73	HACCK29	HADAB60	HADAM31	HADCL19	HADCZ65	HADDC04	HADDP23	HADDP51	HADDR24
Gene No.	20	21	22	23	24	25	26	27	28	29	30

Last AA of ORF	31	15	32	32	46	99	38	57	55	27	36
First AA of Secreted Portion	19		19	19	29	20	31	20	26	21	61
Last AA of Sig Pep	18		18	18	28	16	30	61	25	20	18
First AA of Sig Pep	_	1	_	1	1	1	1	1	1	1	1
AA SEQ ID NO: Y	2638	2639	2640	2641	2642	2643	2644	2645	2646	2647	2648
	154	201	169	169	94	377	212	24	147	223	187
5' NT of of of Start AA of Codon Signal Pep	154	201	169	169	94	377	212	24	147	223	187
	830	4054	452	625	1193	1594	1762	1042	852	1120	1278
5' NT 3' NT of Ol Clone Clone Seq.	1	1	1	-		1	1	1	15	1	1
Total NT Seq.	830	4054	452	625	1193	1594	1762	1042	855	1120	1278
NT SEQ D NO:	41	42	43	44	45	46	47	48	49	50	51
Vector	pSport1	pSport!	pSport1	pSport1	pSport1						
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HADET62	HADEY08	HADEY22	HADEY22	HADFB84	HADFD01	HADFD10	HADFK11	HADFT44	HADFW20	HADFX10
Gene No.	31	32	33	34	35	36	37	38	39	40	41

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Last AA of	ORF	34	45	23	123	123	87	31	41	35	17	58
	Portion	24	26	15	49	46	51	18	36	23	11	27
	Рер	23	25	14	48	48	20	17	35	22	10	26
First AA of Sig	Pep	1	1	1	1	1	1	1	1	1	1	1
' 02 ~	Y	2649	2650	2651	2652	2653	2654	2655	2656	2657	2658	2659
	Signal Pep	83	47	992	1327	731	186	542	82	32	205	106
5' NT of Start	Codon Signal Pep	83	47		1327	731	186	542	82	32		901
3' NT of Clone Seq.		742	899	1913	1927	1331	1733	2722	1094	1839	1964	1330
S' NT 3' NT of Of Clone Clone Seq. Seq.		1	1	763	1149	553	1	364	1	П	-	1
	Seq.	742	1033	1913	1992	1386	1733	2722	1094	1839	1964	1330
SEQ NO:	X	52	53	54	55	99	57	58	59	09	61	62
1	Vector	pSport1	pSport1	pBluescript	pSport1	pSport1	pBluescript SK-	pBluescript SK-	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z	and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203960 04/26/99	203960 04/26/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99	203917 04/08/99
	Clone ID	HADFY80	HADGD93	HADMA77	HADXA10	HADXA10	HAFBB15	HAFBL14	HAGAB62	HAGAB83	HAGAE84	HAGAF75
Gene	No.	42	43	4	45	46	47	48	49	20	51	52

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Last AA of ORF	19	32	9	18	30	32	32	37	36	25	17
First AA of Secreted Portion	16	21	16	16	28	29	19	30	25	22	
Last AA of Sig Pep	15	20	15	15	27	28	18	29	24	21	
AA First SEQ AA ID of NO: Sig Y Pep	1		1	-	1	1	1	1	1	-	1
AA SEQ ID Y	2660	2661	2992	2663	2664	2665	3666	2667	2668	2669	2670
5' NT of First AA of Signal Pen		285	128	358	27	59	54	132	86	86	725
5' NT of Start Codon		285	128	358	27	59	54	132	86	86	725
3' NT of Clone Seq.	1504	1828	1280	1528	1458	1538	557	1568	1228	1715	1846
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-	-							-		456
Total NT Seq.	1504	1828	1280	1528	1458	1538	557	1568	1228	1715	1896
NA NO:	63	64	92	99	<i>L</i> 9	89	69	70	71	72	73
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203918	203917 04/08/99	203917 04/08/99	203917	203979 04/29/99	203917 04/08/99	203917 04/08/99	203918	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HAGAK40	HAGAU43	HAGAZ36	HAGBC57	HAGBL31	HAGBO09	HAGB012	HAGB051	HAGBS89	HAGBV06	HAGBV25
Gene No.	53	54	55	26	57	58	59	09	61	62	63

Last AA of ORF	59	6	41	99	39	35	30	36	33	3	33
First AA of Secreted Portion	22		31	18	L7	19	17	<i>L</i> 1	32		19
Last AA of Sig Pep	21		30	<i>L</i> 1	26	18	16	16	31		18
First Last AA AA of of Sig Sig Pep	1	1	1	1	1	-	-	1	-	-	1
	2671	2672	2673	2674	2675	2676	2677	2678	2679	2680	2681
	91	605	150	31	42	91	744	297	45	110	130
5' NT of of of Start AA of Codon Signal Pep	16	509	150		42	91	744	297	45	110	130
3' NT of Clone Seq.	2075	1592	1324	1214	1338	1686	1634	2012	1322	941	1874
5' NT 3' NT of of Clone Clone Seq. Seq.	8	479	1	-	-	54	277	1	1	1	2
Total NT Seq.	2075	1592	1324	1214	1338	1686	1634	2012	1322	941	1874
NT SEQ ID NO:	74	75	9/	11	78	62	80	81	82	83	84
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR						
ATCC Deposit No.Z and Date	203917 04/08/99	203917	203917 04/08/99	203917	203917 04/08/99						
cDNA Clone ID	HAGBV29	HAGCC87	HAGCH67	HAGCI69	HAGCT33	HAGCZ70	HAGDC73	HAGDG84	HAGDH85	HAGDI69	HAGDJ53
Gene No.	64	99	99	<i>L</i> 9	89	69	70	71	72	73	74

Last AA of ORF	34	33	180	35	39	2	39	31	35	17	37
First AA of Secreted Portion	25	22	28	23	19		24	25	19		17
	24	21	27	22	18		23	24	18		16
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	—	-	-	_	-	-
AA SEQ ID NO: Y	2682	2683	2684	2685	2686	2687	2688	2689	2690	2691	2692
	81	218	338	198	06	223	932	165	120	162	226
5' NT of of of Start AA of Codon Signal Pep	81	218	338	198	90	223	932	165	120	162	226
	643	1669	945	2007	1687	952	1388	1759	810	1675	1205
s' NT of Clone Seq.	1	1	144	-	-	П	744	1	1	1	-
Total NT Seq.	643	6991	948	2007	1687	952	1410	1759	810	1675	1205
NT SEQ ID NO:	85	98	87	88	68	06	91	92	93	94	95
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917	203917	203917 04/08/99	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HAGDJ56	HAGDL51	HAGDO70	HAGDT30	HAGDW68	HAGDX84	HAGEK37	HAGEK86	HAGEP30	HAGEQ58	HAGEQ67
Gene No.	75	9/	77	78	79	80	81	82	83	8	85

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Last AA of ORF	53	31	16	16	61	5	92	=	120	44
First AA of Secreted Portion	21	25			17		27		23	22
	20	24			16		76		22	21
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	-	-	1	1	1	1	-	1		
AA SEQ ID NO: Y	2693	2694	2695	2696	2697	2698	2699	2700	2701	2702
	175	18	909	606	13	266	208	243	830	105
5' NT of of of Start AA of Codon Signal Pep	175	18	606	909	13	266	208	243		105
	484	1069	1434	1434	1067	1844	2122	931	1683	1270
5' NT 3' NT of of Clone Clone Seq. Seq.	1	1	373	373	1	-		-	633	-
Total NT Seq.	484	1069	1475	1475	1067	1844	2122	931	1683	1270
SEQ SEQ NO:	96	76	86	66	100	101	102	103	104	105
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917	203917	203917 04/08/99	203979 04/29/99	203917 04/08/99	PTA- 181 06/07/99	203917 04/08/99	203979	203917 04/08/99
cDNA Clone ID	HAGEU26	HAGEW83	HAGEX49	HAGEX49	HAGFD75	HAGFF43	HAGFJ67	HAGFM58	HAGFT48	HAGFU31
Gene No.	98	87	88	68	8	91	92	93	94	95

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Last AA of ORF	34	5	32	23	39	1	49	72	35	34
First AA of Secreted Portion	21		18	23	32		23	19	26	18
Last AA of Sig Pep	20		17	22	31		22	18	25	17
First AA of Sig Pep	1	1	1	1	1	1	1	-	_	-
AA SEQ Y Ö.	2703	2704	2705	2706	2707	2708	2709	2710	2711	2712
r. 44 –	rep 183	342	28	65	185	595	85	137	286	06
5' NT of Start Codon	183		28	65	185	595	85	137	286	06
3' NT of Clone Seq.	911	1697	1142	926	658	1588	593	1903	1437	2050
5' NT of Clone Seq.		-	-		1	425	1	1	-	1
Total NT Seq.	911	1697	1142	926	859	1588	593	2355	1437	2050
NT SEQ ID NO:	106	107	108	109	110	111	112	113	114	115
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript	pBluescript	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917	203917	203917 04/08/99	203959 04/26/99	203917 04/08/99	203979 04/29/99	203979 04/29/99	PTA- 791 09/27/99	203917 04/08/99	203979 04/29/99
cDNA Clone ID	HAGFW13	HAGHE85	HAGHR18	HAGIB90	HAHEM51	HAHSA76	HAHSD51	HAIBR76	HAIBT20	HAIBV91
Gene No.	96	26	86	66	100	101	102	103	104	105

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Last AA of ORF	5	39	26	25	92	35	72	37	17	99	39
First AA of Secreted Portion		24	24	16	15	20	30	26		22	18
Last AA of Sig Pep		23	23	15	14	61	29	25		21	17
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	-	1	1	-	1	1	-	-		1
	2713	2714	2715	2716	2717	2718	2719	2720	2721	2722	2723
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	363	140	232	255	126	300	282	06	644	661	198
5' NT of Start Codon	363	140	232	255	126	300	282	06	644	661	198
	1361	754	1324	1182	891	1099	1379	581	1259	431	928
5' NT 3' NT of of Clone Clone Seq. Seq.	38	1		1	213	267	1	1	323	П	-
Total NT Seq.	1968	754	1324	1182	911	1099	1379	581	1284	431	876
SEQ BD NO:	116	117	118	119	120	121	122	123	124	125	126
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0 122	pCMVSport 3.0	pCMVSport 3.0	pSport1	pSport1
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979	203918 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99
cDNA Clone ID	HAICE62	HAICL90	HAICV44	HAIDP45	HAJAB88	HAJAZ56	HAMFC67	HAMFQ38	HAMGG01	HANGB24	HANKC93
Gene No.	106	107	108	109	110	111	112	113	114	115	116

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Last AA of ORF	34	52	45	35	35	70	102	36	4	2	16
First Last AA of AA Secreted of Portion ORF	18	23	20	24	24	37	42	21	41		
	17	22	19	23	23	36	41	20	40		
First AA of Sig Pep	1	-	1	-	1	1	1	_		-	-
	2724	2725	2726	2727	2728	2729	2730	2731	2732	2733	2734
5' NT of First AA of Signal Pep	50	125	247	403	403	1872	701	205	33	99	286
5' NT of Start Codon	50	125	247	403	403	1872	701	205		99	
	2157	1585	792	1351	1348	3376	2239	1134	1509	1365	1857
S' NT 3' NT of Clone Seq.	1	1	1	255	255	1593	433	1	1	35	1
Total NT Seq.	2157	1585	792	1351	1371	3397	2812	1145	1509	1365	1857
SEQ NO:	127	128	129	130	131	132	133	134	135	136	137
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203917	203917 04/08/99	203917	203917 04/08/99	203979 04/29/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99	203917 04/08/99
cDNA Clone ID	HAPAD35	HAPBR13	HAPBU09	HAPBU86	HAPBU86	HAPNJ33	HAPNL62	HAPNO50	HAPNY10	HAPPW83	HAPQJ73
Gene No.	117	118	119	120	121	122	123	124	125	126	127

NT of of 5' NT of SEQ AA AA SEQ OF SEQ NT OF SEQ OF	R 138 1810 1 1810 300 2735 1	R 139 1879 1264 1807 1324 1324 2736 1	R 140 1879 1264 1807 1324 1324 2737 1	R 141 556 1 556 20 20 2738 1	R 142 1632 285 1623 515 515 2739 1	R 143 1380 1 1380 172 172 2740 1	R 144 1380 1 1380 251 251 2741 1	R 145 1048 1 1031 170 2742 1	R 146 1882 1 1882 192 192 2743 1	R 147 2254 521 2121 671 2744 1	R 148 284 1 284 79 79 2745 1
ATCC Deposit No.Z and Date Vector	203917 Uni-ZAP XR 04/08/99	203917 Uni-ZAP XR 04/08/99	203917 Uni-ZAP XR 04/08/99	203917 Uni-ZAP XR 04/08/99	203918 Uni-ZAP XR 04/08/99	203917 Uni-ZAP XR 04/08/99	203979 Uni-ZAP XR 04/29/99	203917 Uni-ZAP XR 04/08/99	203917 Uni-ZAP XR 04/08/99	203959 Uni-ZAP XR 04/26/99	203917 Uni-ZAP XR
AT CDNA No Clone ID and	HAPQK26 203	HAPQU71 203	HAPQU71 203 04/0	HAPQW18 203 04/0	HAPQX44 203 04/0	HAPRK55 203 04/0	HAPSH37 203 04/2	HAQBG57 203 04/0	HAQBY85 203 04/0	HAQBZ15 203 04/2	HAQCE18 203
			1	 	 	+	 	135	136	 	138

Last AA of ORF	25	97	334	33	20	38	5	30	29	45	38
First I AA of AP Of Portion C		31	22	19	26	31		28	23		24
		30	21	18	25	30		27	22		.23
First Last AA AA of of Sig Sig Pep	1		1	1	1	1	1	1	1	1	1
	2746	2747	2748	2749	2750	2751	2752	2753	2754	2755	2756
	68	225	422	435	1767	136	669	92	154	134	26
5' NT of of of Start AA of Codon Signal Pep	68	225	422	435	1767	136		9/	154	134	26
3' NT of Clone Seq.	1495	1245	1961	936	2090	1357	810	811	1010	1199	434
5' NT 3' NT of of Clone Seq. Seq.	7	1	189	242	1557		-	-	-		-
Total NT Seq.	1615	1245	1961	936	3853	1357	810	811	1010	1199	434
SEQ SEQ NO:	149	150	151	152	153	154	155	156	157	158	159
Vector	Uni-ZAP XR	pBluescript SK-	pBluescript SK-	pBluescript SK-	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203918 04/08/99	203979	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99	203917 04/08/99
cDNA Clone ID	НАОСР94	HARAE26	HARAT69	HARAZ81	HASAU26	HASAX57	HASAY07	HATAE01	HATAG52	HATAL05	HATBA90
Gene No.	139	140	141	142	143	144	145	146	147	148	149

F St	~	2		<u>, </u>	0	2	46		2	33
Last AA of ORF	38	52	31	15	9	45	4	31		3
First AA of Secreted Portion	37	20	19		17	11	19	16		
Last AA of Sig Pep	36	19	18		16	10	18	15		
	_	-	-	1	1	1	1	1	-	
AA SEQ ID NO: Y	2757	2758	2759	2760	2761	2762	2763	1066 2764	2765	2766
	196	620		237	259	196	428	1066	86	50
5' NT of of of Start AA of Codon Signal Pep	196	620	158	_	259		428	1066		20
3' NT of Clone Seq.	716	2485	1108	930	794	1145	1276	1292	1340	2097
5' NT 3' NT of of Clone Seq.	1	506	1	1	1	,	282	729	Н	
Total NT Seq.	716	2503	1108	930	794	1145	1927	1316	1340	2097
SEQ NO:	160	161	162	163	164	165	166	167	168	169
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203918 04/08/99	203979 04/29/99	203979 04/29/99	203979	PTA- 794 09/27/99	203917	203917 04/08/99	203917 04/08/99	203979 04/29/99
cDNA Clone ID	HATBM71	HATCF80	HATCI67	HATCJ27	HATCS79	HATCX03	HATDE03	HATDF41	HATDH23	HATDH55
Gene No.	150	151	152	153	154	155	156	157	158	159

Last AA of ORF	32	70	46	37	28	32		82	39	45
	8	7	4	3	7	_ω	4		01	7
First AA of Secreted Portion	23	17	14	25		21	34	32	26	17
Last AA of Sig Pep	22	16	13	24		70	33	31	25	16
First AA of Sig Pep	1	1	1		-			-	-	-
AA SEQ ID NO: Y	2767	2768	2769	2770	2771	2772	2773	2774	2775	2776
5' NT of of First Start AA of Codon Signal Pep	75	23	232	139	154	230	2203	265	140	228
5' NT of Start Codon	75	23	232	139	154		2203		140	228
	958	1302	558	1679	1335	263	2394	1279	1009	561
5' NT 3' NT of of Clone Clone NT Seq. Seq.	_	1	-	1	1	1	2100	25	П	
Total NT Seq.	958	1302	558	1679	1335	563	176 2418	1308	1009	561
SEQ NS DS:	170	171	172	173	174	175	176	177	178	179
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript SK-	pBluescript SK-
ATCC Deposit No.Z and Date	203917	203979 04/29/99	203917	203917	203917	203917 04/08/99	PTA- 181	203979	PTA- 791	148
cDNA Clone ID	HATDO84	HATDU01	HATDW05	HATEF13	НАТЕF64	НАТЕН40	HATE122	HAUCC84	HAWAS41	HAWBA65
Gene No.	160	161	162	163	164	165	166	167	168	169

F A St			_	10	41	11	63	29	10	39
Last AA of ORF		30	47		4		9	<u>ν</u>	_	-3
First AA of Secreted Portion		23	24	6	15		18	12		18
Last AA of Sig Pep		22	23	∞	14		17	=		17
First AA of Sig Pep	1	-	1	1	1	-	-	-	1	-
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	<i>TTT</i> 2	2778	2779	2780	2781	2782	2783	2784	2785	2786
5' NT of SEQ A of First ID of Start AA of NO: Start AA of NO: Scoon Signal Y Pep	645	17	101	168	646	805	106	199	96	124
5' NT of Start Codon		17	101				106	199	96	124
3' NT of Clone Seq.	1718	422	2234	307	1758	1056	470	932	1953	1008
S' NT 3' NT of of Of Clone Clone NT Seq. Seq.	-	_	-	1	585	1	1	-	-	1
Total NT Seq.	1718	422	2234	307	1758	1056	470	932	1953	1008
SEQ BD NO:	180	181	182	183	184	185	186	187	188	189
Vector	pSport1	pSport1	pSport1	pSport1	pCMVSport 1	pCMVSport 1	pCMVSport 1	pCMVSport 1	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	PTA- 181 06/07/99	PTA- 791 09/27/99	203917	203917	203917	203917 04/08/99	203917 04/08/99	203917	203917 04/08/99
cDNA Clone ID	HBAGH64	HBAGV01	HBAMC50	HBAMC57	HBBBA42	HBBBB08	HBBBE83	HBBMA11	HBCAK10	HBCAK80
Gene No.	170	171	172	173	174	175	176	177	178	179

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Last AA of ORF	34	31	46	36	55	1		37	113	38	162
First AA of Secreted Portion	31	23	22	. 25	19			22	28	25	59
Last AA of Sig Pep	30	22	21	24	18			21	27	24	28
First AA of Sig Pep	1	1	1				-	-	-	-	-
AA SEQ D NO:	2787	2788	2789	2790	2791		2792	2793	2794	2795	2796
5' NT of First AA of Signal	Pep 320	281	225	298	161	- 1	109	98	06	163	53
5' NT of Start Codor	320	281	225	298	161		109	98		163	53
3' NT of Clone Seq.	421	1086	1038	718	646		893	519	453	414	674
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-	1	-	267	_		89	1	1	П	1
Total NT Sea.	421	1086	1038	765	899		893	519	453	415	674
SEQ BD NO:		191	192	193	194		195	196	197	198	199
Vector	Uni-ZAP XR	Uni-ZAP XR	pSport1	Uni-ZAP XR	Uni-ZAP XR		Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z	203917	203917	203917	203917 04/08/99	PTA- 794	09/27/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA	∞	HBCAY17	HBCGE46	HBGBA14	HBGBE75		HBGBP22	НВ GFQ34	HBGML95	HBGMT60	HBHAA53
Gene	180	181	182	183	184		185	186	187	188	189

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Last AA of ORF	18	30	71	183	24	38	35	15	2	23
First AA of Secreted Portion	16	19	31	20	6	23	29	14		
Last AA of Sig Pep	15	18	30	19	8	22	28	13	!	
AA First Last SEQ AA AA ID of of of NO: Sig Sig Y Pep Pep	-	1	1	1	1	1	1	1	1	1
AA SEQ ID NO: Y	2797	2798	2799	2800	2801	2802	2803	2804	2805	2806
5' NT of First AA of Signal		17	153	929	2	71	112	27	235	234
5' NT of Start Codon	202	17		655		71				234
3' NT of Clone Seq.	1246	1510	1259	2089	725	926	1248	824	2206	1421
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-	-	-	591	-	1	1	-	1	1
	1246	1510	1259	2101	725	926	1248	824	2206	1421
SEQ BD NO:	200	201	202	203	204	205	206	207	208	209
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959	203917 04/08/99	PTA- 794 09/27/99	203979 04/29/99	203979 04/29/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HBIAU43	HBIAW58	HBIBB20	HBIBF26	HBIBM33	HBIBN67	HBIBQ69	HBIBR38	HBIBR61	HBIBS33
Gene No.	190	191	192	193	194	195	196	197	198	199

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Last AA of ORF	33	18	20	13	4	39	8	40	62	36	33
First AA of Secreted Portion	. 22		19			26			23	61	16
Last AA of Sig Pep	21		18			25			22	18	15
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	1	1	1
AA SEQ D NO: Y	2807	2808	2809	2810	2811	2812	2813	2814	2815	2816	2817
5' NT of First AA of Signal Pep	65	146	190	329	274	06	173	50	146	55	369
5' NT of Start Codon	65		190		274	06	173	20	146	55	369
5' NT 3' NT of of Clone Clone Seq.	989	1408	282	1729	781	2115	1148	1131	1117	935	1997
S' NT 3' NT of Clone Clone Seq. Seq.	1	1	1	184	1	-	-	1	-	-	1
Total NT Seq.	630	1408	785	1767	781	2115	1148	1131	1117	963	2884
NT SEQ D NO:	210	211	212	213	214	215	216	217	218	219	220
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99	203979 04/29/99
cDNA Clone ID	HBIBT13	HBIBZ20	HBICB80	HBJAC40	HBJAV56	HBJAY14	HBJBQ69	HBJBR40	HBJCH46	HBJCR17	HBJCS26
Gene No.	200	201	202	203	204	205	206	207	708	209	210

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Last AA of ORF	26	6	17	25	18	45	7	14	59	59
First AA of Secreted Portion	24		17	24		22	,	11	21	21
	23		16	23		21		10	20	20
First AA of Sig Pep	1	-	1		1	1	1	1	1	-
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	2818	2819	2820	2821	2822	2823	2824	2825	2826	5195
5' NT AA of SEQ First ID AA of NO: Signal Y		173	265	141	99	48	1663	264	180	851
5' NT of Start Codon	323	173			95	48	1663		180	851
3' NT of Clone Seq.	1014	743	1118	1485	1249	2082	2294	1254	895	1552
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	-	1	1	1	—	-	1580	1	-	672
Total NT Seq.	1014	743	1118	1485	1249	2082	2294	1255	895	2598 2178
SEQ BD NO:	221	222	223	224	225	226	227	228	229	2598
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917	203917	PTA- 181 06/07/99	203959	203979 04/29/99	PTA- 792 09/27/99	PTA- 792 09/27/99
cDNA Clone ID	HBJCW24	HBJDC57	HBJDR18	HBJDR83	HBJEE51	HBJEL21	HBJFH84	HBJFJ14	HBJFJ26	HBJFJ26
Gene No.	211	212	213	214	215	216	217	218	219	219

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Last AA of ORF	34	34	38	31	77	13	<i>L</i> 9	18	21	47	35
First AA of Secreted Portion	31	31	28	28	23		35		21	22	15
	30	30	27	27	22		34		8	21	14
First Last AA AA of of Sig Sig Pep	1	-	-	1	1	_	1	-	-	-	-
AA SEQ ID NO: Y	2827	2828	2829	2830	2831	2832	2833	2834	2835	2836	2837
	246	246	423	354	14	181	189	121	200	193	443
5' NT of of of Start AA of Codon Signal Pep	246	246	423	354		181	189	121	200	193	443
3' NT of Clone Seq.	1208	1165	1021	1661	477	779	972	1885	1251	1252	1256
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	335	190	-	1	1	64	1	1	224
Total NT Seq.	1208	1165	1021	1661	477	779	972	1885	1251	1252	1256
NT SEQ BD NO:	230	231	232	233	234	235	236	237	238	239	240
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HBJFJ83	HBJFJ83	HBJFP47	HBJFR77	HBJFU30	HBJFX41	НВЈНО83	HBJHS92	HBJHT01	HBJHT01	HBJHW06
Gene No.	220	221	222	223	224	225	226	227	228	229	230

Last AA of	19	29	36	18	348	34	29	58	82	123	72
First AA of Secreted Portion (13	16	21		19	29	18	31	15	22	27
Last AA of Sig Pep	12	15	70		18	28	17	30	14	21	26
First AA of Sig Pep	-	-		1	1	1	1	-	_	1	-
AA SEQ ID NO: Y	2838	2839	2840	2841	2842	2843	2844	2845	2846	2847	2848
	1800 2838	236	228	284	309	20	222	26	47	441	413
5' NT of of of Start AA of Codon Signal Pep	1800	236	228	284		20		99			413
	2809	1257	724	1099	1701	1120	1832	1247	621	998	2779
5' NT 3' NT of of Clone Seq. Seq.	1522	1	1	128	86	-	1	1	1	I	242
Total NT Seq.	2858	1363	724	1099	1703	1120	1832	1247	621	998	3057
X SEQ	241	242	243	244	245	246	247	248	249	250	251
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1	pSport1	pSport1	pSport1	pBluescript
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99
cDNA Clone ID	HBJIR14	HBJJA26	HBJJX02	нвл.н78	HBJND04	HBJND57	HBKDF66	HBKEA94	HBKEE60	HBKEI41	HBMBD51
Gene No.	231	232	233	234	235	236	237	238	239	240	241

Last AA of ORF	13	37	37	31	70	50	20	82	156	31
First AA of Secreted Portion (23	17	16			20	29	23
			22	16	15			61	28	22
First AA of Sig Pep	1	1	-	-	1	1	1	1	-	1
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	2849	2850	2851	2852	2853	2854	2855	2856	2857	2858
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	4310 2849	45	31	69	100	1815 2854	1815 2855	184	202	193
5' NT of Start Codon		45	31	69	100			184	202	193
	5091	931	1162	807	2284	3480	3480	711	692	973
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	4063	1	1	1	2	1563	1563	1	629	1
Total NT Seq.	5279	931	1162	807	2284	4043	4044	711	1113	985
SEQ NO:	252	253	254	255	256	257	258	259	260	261
Vector	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917	203917	203917 04/08/99	203917 04/08/99		PTA- 181 06/07/99	PTA- 181 06/07/99	203917 04/08/99	203979 04/29/99	203917 04/08/99
cDNA Clone ID	HBMBD73	HBMBE33	HBMBM17	HBMCL59	НВМСМ96	HBMCQ74	HBMCQ74	HBMCT40	HBMDM08	HBMSN62
Gene No.	242	243	244	245	246	247	248	249	250	251

		ATCC		NT		5' NT 3' NT of	3' NT of	5' NT	5° NT of	AA SEQ	First AA	Last AA	First	Last
Gene	cDNA	Deposit No.Z		ВÃ	Total NT	Total Clone Clone NT Seq. Seq.	Clone Seq.	of Start	First ID AA of NO:	ΑÖ	of Sig	of Sig	AA of Secreted	AA of
No.	_	and Date	Vector	×	Seq.			Codon Signal Pep	Signal Pep	Y	Pep	Pep	Portion	OKT
252	HBMSO30	203959	Uni-ZAP XR	262	778	1	778	172	172	2859	_	16	17	17
253	HBMTM50		Uni-ZAP XR	263	1234	-	1234	4	4	2860	-			4
254	HBMUD59		Uni-ZAP XR	264	876	-	876	270	270	2861	-	21	22	126
255	HBMUI10	203917	Uni-ZAP XR	265	822	1	822	249	249	2862	1	28	29	39
256	HBMUJ48	203917	Uni-ZAP XR	266	513	1	513	180	180	2863	-	17	18	24
257	HBMUR39	203917	Uni-ZAP XR	267	888	1	888	135	135	135 2864	-	18	19	23
258	HBMVF65	203917 04/08/99	Uni-ZAP XR	268	1064	175	1063		179	2865	-	9	7.1	38
259	HBMVF65	203917 04/08/99	Uni-ZAP XR	269	1282	378	1265	382	382	2866	-	19	17	85
260	HBMWC39	203917 04/08/99	Uni-ZAP XR	270	1154	-	1154		112	2867	_	9		18
261	HBMWJ92	203917 04/08/99	Uni-ZAP XR	271	1810	1	1810	82	82	2868	-	17	18	40

Last AA of ORF	35	=	24	26	37	39		148	7	10
	<u>~</u>			.,	(,,					
First AA of Secreted Portion	17		14	21	31	34		36		
Last AA of Sig Pep	16		13	20	30	33		35		
First AA of Sig Pep	1	1	1	1	1	1	-	-	-	П
AA SEQ ID NO: Y	2869	2870	2871	2872	2873	2874	2875	2876	2877	2878
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	19	301	261	71	20	16	203	2711	519	338
5' NT of Start Codon	19	301		71	70	16		2711	519	338
3' NT of Clone Seq.	1345	1831	1139	618	1119	1233	426	3231	894	1599
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-	906	1	-	1	1	-	2578	295	1
Total NT Seq.	1345	1831	1139	618	1121	1233	426	3244	894	1778
X S B S X	272	273	274	275	276	277	278	279	280	281
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1
ATCC Deposit No.Z and Date	PTA- 791 09/27/99	203917	203917	203917	203979	203917 04/08/99	203917	203979	203917 04/08/99	203979 04/29/99
cDNA Clone ID	HBMWS52	HBMXE34	HBMXG01	HBMXG76	HBMXM05	HBMXW83	HBNAE74	HBNAX16	HBNAZ35	HBODK40
Gene No.	262	263	264	265	266	267	268	269	270	271

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Last AA of ORF	∞	15	18	7	LL	37	62	62	42	48
First AA of Secreted Portion		15			30		21	21	22	23
Last AA of Sig Pep		14			56		20	20	21	22
AA First Last SEQ AA AA Of Of Of Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	1	1
AA SEQ ID NO: Y	2879	2880	2881	2882	2883	2884	2885	2886	2887	2888
5' NT of First AA of Signal Pep	303	74	119	292	741	208	1256 2885	183	833	182
5' NT of of of Start AA of Codon Signal Pep	303	74	119		741	208			833	182
3' NT of Clone Seq.	2498	195	1262	1371	2345	1015	1587	540	1494	1504
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	1	П	1	1079		710	1
Total NT Seq.	2498	195	1262	1371	2345	1015	1708	540	1494	1504
NT SEQ D NO:	282	283	284	285	286	287	288	289	290	291
Vector	pSport1	Uni-ZAP XR	Uni-ZAP XR	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	ZAP Express
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203917 04/08/99	203979 04/29/99	203917 04/08/99		203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HBODV76	HBPAD89	HBPAF39	HBQAC45	HBQAC72	HBQAE37	HBSAJ63	HBSAJ63	HBSDD24	HBWBD25
Gene No.	272	273	274	275	276	277	278	279	280	281

Last AA of ORF	=	40	36	28	46	119	65	4	17	37
First AA of Secreted Portion		18	31	28	31	22	23		14	15
		17	30	27	30	21	22		13	14
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	-		-	-	_	1	-	1	-	1
4	2889	2890	2891	2892	2893	2894	2895	2896	2897	2898
	215	1097	152	87	85	2163 2894	165	349	260	366
5' NT of of of Start AA of Codon Signal Pep	215		152	87		2163	165	349	260	366
	1759	2385	1377	2043	713	2791	2017	1273	1879	2513
5' NT 3' NT of of Clone Seq. Seq.	1	856	1	-	П	2044	1	1	1	355
Total NT Seq.	1759	2406	1377	2043	713	2791	2017	1273	1879	2520
SEQ NO:	292	293	294	295	296	297	298	299	300	301
Vector	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express
ATCC Deposit No.Z and Date	203917	203959	203917	203979 04/29/99	203917 04/08/99	203917 04/08/99	PTA- 181 06/07/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HBXAS93	HBXAT27	HBXAW57	HBXB129	HBXBM24	HBXBM78	HBXCD59	HBXCE43	HBXCG08	HBXCM52
Gene No.	282	283	284	285	286	287	288	289	290	291

Last AA of ORF	55	33	∞ .	∞	24	36	41	33	123	37
First AA of Secreted Portion (31	23			23	26	21	18	22	27
	30	22			22	25	20	17	21	26
First Last AA AA of of Sig Sig Pep	1	1	1	1	_	1	-	1	1	1
AA SEQ ID NO: Y	2899	2900	2901	2902	2903	2904	2905	2906	2907	2908
5' NT of First AA of Signal Pep	231	158	1040	1040 2902	<i>L</i> 9	187	411	15	124	122
5' NT of Start Codon	231	158	1040	1040	29	187	411		124	122
3' NT of Clone Seq.	896	1235	2307	2307	1057	1948	617	1647	869	1467
5' NT 3' NT of of Clone Clone Seq. Seq.	T	1	588	885	-	1	1	1	1	1
Total NT Seq.	896	1235	2311	2311	1057	1948	622	1647	869	1467
NT SEQ ID NO:	302	303	304	305	306	307	308	309	310	311
Vector	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	PTA- 791 09/27/99	PTA- 791 09/27/99	203917 04/08/99	203918 04/08/99	203959 04/26/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	нвхс003	HBXCR15	HBXDL52	HBXDL52	HBXDN08	HBXDN65	HBXFA04	HBXFE64	HBXF133	HBXFP72
Gene No.	292	293	294	295	296	297	298	299	300	301

Last AA of ORF	16	182	120	120	36	15	48	37	40	36
First AA of Secreted Portion		2	33	33	35		22	30	25	31
		1	32	32	34		21	29	24	30
First AA of Sig Pep	1	1	1	1	1	1	1	1	1	1
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	2909	2910	2911	9615	2912	2913	2914	2915	2916	2917
5' NT of First AA of Signal Pep	20	125	52	31	08	268	760	6	92	30
5' NT of Start Codon	20	125			08	268		6	92	30
	1551	1252	2516	2469	2483	1663	1531	223	2015	1964
5' NT of Clone Seq.	1	1	1	1			H	-	_	
Total NT Seq.	1551	1252	2516	2469	2483	1663	1531	223	2015	1964
NT SEQ ID NO: X	312	313	314	2599	315	316	317	318	319	320
Vector	ZAP Express	ZAP Express	pSport1	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR				
ATCC Deposit No.Z and Date	203917 04/08/99	203979 04/29/99	203917 04/08/99	203917 04/08/99	PTA- 1838 05/09/00	203917 04/08/99	203917 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203979 04/29/99
cDNA Clone ID	HBXFS31	HBXFW01	HBXGE12	HBXGE12	HBXGL91	HBXGM24	HBZAI75	HCABP33	HCABW10	HCACZ65
Gene No.	302	303	304	304	305	306	307	308	309	310

Last AA of ORF	39	182	142	35	57	27	40	244	48	71
First AA of Secreted Portion	56	56	56	18	19	27	29	19	16	26
Last AA of Sig Pep	25	25	25	17	18	26	28	18	15	25
First AA of Sig Pep	-		1	1	1	1	1	—		-
AA SEQ D NO: Y	2918	2919	2920	2921	2922	2923	2924	2925	2926	2927
	93	342	427	12	495	126	102	136	260	1669 2927
5' NT of of of Start AA of Codon Signal Pep	93	342	427	12	495	126		136		1669
	1650	191	853	803	999	1454	853	1117	685	2630
5' NT of Clone Seq.	1	288	373	-	267	-			-	1480
Total NT Seq.	1650	924	1015	803	999	1454	853	1117	685	2630
SEQ NO:	321	322	323	324	325	326	327	328	329	330
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203979 04/29/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917	203917 04/08/99	PTA- 794 09/27/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HCBAB34	HCDAA24	HCDAA24	HCDAF17	НСДАН02	HCDAP33	HCDAR40	HCDAS02	HCDBE76	HCDB032
Gene No.	311	312	313	314	315	316	317	318	319	320

Last AA of ORF	4	41	139	2	6	12	84	31	84	59
First L AA of Secreted Portion		∞	36				16	29	32	34
	10	7	35				15	28	31	33
First AA of Sig Pep		-	-	1	1	1	-	1	1	-1
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	2928	2929	2930	2931	2932	2933	2934	2935	2936	2937
5' NT AA of SEQ First D AA of NO: Signal Y Pep	498	565	32	46	216	899	274	75	266	83
5' NT of of of Start AA of Codon Signal Pep			32	46	216	668	274	75		83
	219	858	1538	1085	1046	1202	419	1691	1744	957
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	409	21	1	1		514	1	1	1	1
	229	858	1538	1085	1046	1422	419	1691	1744	957
SEQ NO:	331	332	333	334	335	336	337	338	339	340
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99	203917 04/08/99	203917 04/08/99	PTA- 793 09/27/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HCDBW67	HCDBZ31	HCDCB03	HCDCE51	HCDCI42	HCDDB15	HCDDX81	HCDDY28	HCDEB19	HCDEN46
Gene No.	321	322	323	324	325	326	327	328	329	330

r _T		- 1		— т			I			
Last AA of ORF		4	38	9	133	∞	6	29	90	45
First AA of Secreted Portion		29	38	22	11			22	33	17
		28	37	21	10			21	32	16
First AA of Sig Pep	1	1	-	1	1	-	1	-	1	
AA SEQ ID NO: Y	2938	2939	2940	2941	2942	2943	2944	2945	2946	2947
5' NT of First AA of Signal Pep	271	101	106	23	374	757	279	102	99	142
5' NT of Start Codon		101	106	23	374				99	
3' NT of Clone Seq.	1032	1390	1587	1461	1651	1720	1247	1830	977	1893
S' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1	1	-	-		1	-	1	-	
Total NT Seq.	1032	1390	1590	1461	1651	1720	1247	1830	21.6	1893
NT SEQ ID NO:	341	342	343	344	345	346	347	348	349	350
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203959	203917 04/08/99	203917 04/08/99	203917 04/08/99	PTA- 792 09/27/99	203917	203917 04/08/99
cDNA Clone ID	HCDES69	HCE1D45	HCE1N56	HCE1T53	HCE1Y27	HCE1Y34	HCE2B57	HCE2E47	HCE2I23	НСЕ2Р90
Gene No.	331	332	333	334	335	336	337	338	339	340

Last AA of ORF	45	9	99	87	22	46	51	99	99	99
First AA of Secreted Portion (19		19	34		17	15	33	33	33
	18		18	33		16	14	32	32	32
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	-		1	-	1	1	1	1	-
AA SEQ ID NO: Y	2948	2949	2950	2951	2952	2953	2954	2955	2956	2957
	51	290	682	73	1075		789	1386	1386 2956	1386 2957
5' NT of of of Start AA of Codon Signal Pep	51		682	73	1075	221	789	1386	1386	1386
	847	1230	2575	1100	2087	402	3114	2736	2736	1245 2736
5' NT 3' NT of of Of Clone Clone NT Seq. Seq.	1	-	501	17	836	1	262	1245	1245	1245
Total NT Seq.	847	1230	2575	1100	2129	709	3145	2746	2736	360 2736
X SEQ N	351	352	353	354	355	356	357	358	359	360
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	PTA- 791 09/27/99	203917 04/08/99	203917 04/08/99	203918 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HCE3A54	HCE3C46	HCE3D58	нсезD89	HCE3J43	HCE3L04	HCE3N23	HCE3R01	HCE3R01	HCE3R01
Gene No.	341	342	343	344	345	346	347	348	349	350

Last AA of ORF	22	09	9	9/	32	29	49	39	129	99	55
First AA of Secreted Portion (47	47	34	17	24	29	32	31	29	21
		94	46	33	16	23	28	31	30	28	70
First AA of Sig Pep	-		1	1	1	1	—			1	-
AA SEQ ID NO: Y	2958	2959	2960	2961	2962	2963	2964	2965	2966	2967	2968
	376	1403	814	91	286	204	288	114	197	124	71
5' NT of of of Start AA of Codon Signal Pep	376	1403	814			204	288	114	197	124	71
	2025	2605	2016	840	1597	1714	2328	2207	1069	1436	1419
5' NT 3' NT of Of Clone Seq. Seq.	58	1344	755	-	23		1	1	1	-	
Total NT Seq.	2046	2636	2047	840	4151	1714	2329	2207	1069	1436	1419
SEQ NO:	361	362	363	364	365	366	367	368	369	370	371
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203979 04/29/99	203979 04/29/99	203979	203917	203917	203917	203979 04/29/99	203917 04/08/99	203979 04/29/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HCE3R46	НСЕ4Н32	HCE4H32	HCE4T64	HCE4W88	HCE5B62	НСЕ5Н86	HCE5J64	HCEBF54	HCECO77	НСЕДН42
Gene No.	351	352	353	354	355	356	357	358	359	360	361

Last AA of ORF	46		9	34	21	35	_	28	4	
First AA of Secreted Portion	24		17	12		27		25		
	23		16	11		26		24		
irst AA of of Sig	1		1		1	1	-		-	1
AA SEQ ID NO:	5962	2970	2971	2972	2973	2974	2975	2976	2977	2978
5' NT AA F of SEQ A First D AA of NO: Signal Y Pep	<i>L</i> 9	981	126	809	242	244	276	18	282	223
	<i>L</i> 9		126			244	276	18		223
	2396	1984	1425	1953	966	1165	1381	775	1474	2100
5' NT 3' NT of Clone Clone Seq. Seq.	1	34	-	255		-	1			-
Total NT Seq.	2396	1984	1425	1953	966	1165	1381	775	1474	2100
SEQ NO:	372	373	374	375	376	377	378	379	380	381
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203979 04/29/99	203917 04/08/99	PTA- 794	203917	203917	203917 04/08/99	203917	203917 04/08/99
cDNA Clone ID	HCEDJ05	HCEDJ26	HCEDN07	HCEDO17	HCEEG48	НСЕЕМ33	HCEEP16	HCEER60	HCEFA10	HCEFA50
Gene No.	362	363	364	365	366	367	368	369	370	371

Last AA of ORF	17	38	13	59	029	,	4	17	26	99	61
First L AA of A Secreted Portion O		23		32	21	9	61	16	21	56	25
	∞	22		31	20	,	18	15	8	25	24
First AA of Sig	-	-	-	1	1	1	_	-		-	
	2979	2980	2981	2982	2983		2984	2985	2986	2987	2988
	232	224	166	260	79		113	L	ľ	72	145
S' NT of of of Start AA of Codon Signal Pep	232	224		260			113	1114	203	72	145
	1607	1432	2280	2241	2455		639	2025	1124	1786	1688
5' NT 3' NT of of Clone Clone Seq. Seq.	1	1	1	66			_	626	1	1	-
Total NT Seq.	1607	1432	2280	2261	2455		639	2534	1124	1786	1688
SEQ NO:	382	383	384	385	386		387	388	389	390	391
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR		Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917	203917	203917 04/08/99	203917	PTA-	96/12/60	203917 04/08/99	203979 04/29/99	203979 04/29/99	203979	203918 04/08/99
cDNA Clone ID	HCEFA94	HCEFC27	HCEFG93	НСЕГН31	HCEFK56		HCEFN51	HCEGG08	HCEGH74	HCEGK81	HCEGS49
Gene No.	372	373	374	375	376		377	378	379	380	381

Last AA of ORF	31	51	٥	149	23	28	99	23	35	38	38
First L. AA of A Secreted O Portion O				21 1		19	23				33
				70		18	22	15	31	32	32
First Last AA AA of of Sig Sig Pep	-	-	-		1	1	1		-	1	_
	2989	2990	2991	2992	2993	2994	2995	2996	2997	2998	2999
			29	175	20	108	37	56	263	242	242
5' NT of of of Start AA of Codon Signal Pep	84	174	29	175	20		37		263	242	242
3' NT of Clone Seq.	1487	2803	1380	1110	1305	1962	1880	878	2320	6991	1668
S' NT 3' NT of of Clone Clone Seq.	1	1	-	1	-	1		1	1	1	_
Total NT Seq.	1487	2834	1380	1140	1305	1962	1880	878	2320	1669	1668
SEQ NO:	392	393	394	395	396	397	398	399	400	401	402
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917	203917 04/08/99	203917	203917	203917	203979	203959	203917 04/08/99	203917 04/08/99	203917	203917 04/08/99
cDNA Clone ID	HCEGU75	HCEGY33	HCEHW24	HCEJL08	HCEJP93	HCELB04	HCEMA08	HCENN67	HCENQ22	НСЕОF01	нсеоғ01
Gene No.	382	383	384	385	386	387	388	389	390	391	392

Last AA of ORF	27	32	59	40	2	28	∞	20	40	4
First AA of Secreted Portion	15	30	23	34		21		18	34	
	14	29	22	33		20		17	33	
AA First Last SEQ AA AA ID of of of NO: Sig Sig Y Pep Pep		1	1	1	1	1	1	1	-	1
	3000	3001	3002	3003	3004	3005	3006	3007	3008	3009
	603	18	1563	232	190	228	223	227	114	223
S' NT of of of Start AA of Codon Signal Pep	603	18	1563	232		228	223	227	114	223
3' NT of Clone Seq.	1677	992	2150	939	641	883	1350	2541	647	635
5' NT 3' NT of clone Clone Seq.	477	1	1376	П	-	1	1	201	Ţ	210
Total NT Seq.	1677	992	2150	939	641	883	1350	2541	647	1203
SEQ NO:	403	404	405	406	407	408	409	410	411	412
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript	pBluescript	pBluescript	pBluescript	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203979 04/29/99	PTA- 181 06/07/99	203917 04/08/99	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HCEON94	НСЕОО67	HCEOV48	HCEPC90	HCEPO08	HCESB03	HCESK44	HCETE08	HCETL19	HCEWD90
Gene No.	393	394	395	396	397	398	399	400	401	402

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Last AA of ORF	38	1	65	40	15	39	3	36	295	36
First AA of Secreted Portion	21		36	28		23		32	34	29
Last AA of Sig Pep	20		35	27		22		31	33	28
AA First SEQ AA ID of NO: Sig Y Pep	1	1	1	1	1	1	1	1	1	1
AA SEQ UO: Y	3010	3011	3012	3013	3014	3015	3016	3017	3018	3019
5' NT of First AA of Signal Pep		141	147	119	120	93	419	65	1854 3018	390
5' NT of Start Codon	158		147	119	120	93		76	1854	390
3' NT of Clone Seq.	1561	2071	066	1780	698	929	1759	1718	2921	1677
S' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	-	1		П	_	1	068	Π	1720	1
Total NT Seq.	1561	2071	066	1780	698	929	1759	1718	2927	1677
SEQ BD NO:	413	414	415	416	417	418	419	420	421	422
Vector	Uni-ZAP XR	Uni-ZAP XR	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203959 04/26/99	203917 04/08/99	PTA- 794 09/27/99	203979 04/29/99
cDNA Clone ID	HCEWE62	HCEZW14	HCFAT42	HCFAT66	HCFBA30	HCFBM77	HCFBV39	HCFCB72	HCFCG91	HCFCM81
Gene No.	403	404	405	406	407	408	409	410	411	412

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Last AA of ORF	37	34	34	9	16	319	101	30	9	49	∞
First AA of Secreted Portion	. 15	28	28	9	14	33	. 16	70	29	37	
Last AA of Sig Pep	14	27	27	5	13	32	15	19	28	36	
First AA of Sig Pep	1	-	1	1	1	1	1		—	-	-
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	3020	3021	3022	3023	3024	3025	5197	3026	3027	3028	3029
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	32	53	191	52	621	121	437	78	44	167	385
5' NT of Start Codon	32	53	161			121		78		167	385
3' NT of Clone Seq.	1343	1523	1691	870	1591	1482	1443	1041	1783	2208	845
of Of Clone Seq.	-	-	57	168	-	-	1	1	1	119	27
Total NT Seq.	1343	1523	1691	870	1622	1482	1464	1041	1783	2208	1097
NT SEQ ID NO:	423	424	425	426	427	428	2600	429	430	431	432
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203979	203917 04/08/99	203917	203917 04/08/99	203917 04/08/99	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HCFCW39	HCFCY49	HCFDD18	HCFLB10	HCFLC03	HCFLJ52	HCFLJ52	HCFLL33	HCFLP48	HCFLQ12	HCFLY20
Gene No.	413	414	415	416	417	418	418	419	420	421	422

Last AA of ORF	158	45	22	94	32	28	49	31	25	46
	1									
First AA of Secreted Portion	2	22	11	28	24	28	20	23		31
Last AA of Sig Pep	1	21	10	27	23	27	19	22		30
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep		-	1	-	-	-	-		-	1
AA SEQ ID NO: Y	3030	3031	3032	3033	3034	3035	3036	3037	3038	3039
5' NT of First AA of Signal Pen	95	217	321	110	112	354	191	448	335	183
5' NT AA F 5' NT of SEQ of First ID Start AA of NO: Codon Signal Y I	95	217		110	112	354	191	448		183
5' NT 3' NT of of Of Clone Clone NT Seq. Seq.	857	382	750	1238	829	802	1148	1095	1393	1597
5' NT 3' NT of of Clone Clone Seq. Seq.	38	1	1	1	1	1	-	1	1	1
	1123	382	750	1238	829	802	1148	1095	1393	1597
NT SEQ ID NO:	433	434	435	436	437	438	439	440	441	442
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	PTA- 794 09/27/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HCFLY20	HCFMA39	HCFMJ40	HCFML07	HCFMR75	HCFMX16	HCFMX88	HCFNM40	HCFNM50	HCFNN16
Gene No.	423	424	425	426	427	428	429	430	431	432

Last AA of ORF	49	34	31	32	216	36	59	55	347	28
First AA of Secreted Portion	20	29	24	27	34	12	19	27	49	
Last AA of Sig Pep	19	28	23	26	33	11	18	26	48	
AA First SEQ AA ID of NO: Sig Y Pep	1	1	1	1	1	1	1	1	1	1
AA SEQ ID NO: Y	3040	3041	3042	3043	3044	3045	3046	3047	3048	3049
5' NT AA of SEQ First ID AA of NO: Signal Y	241	27	3	83	201	234	267	276	168	341
5' NT of Start Codon	241	27	3	83	201	234	267	576	168	
3' NT of Clone Seq.	1641	1470	604	981	1653	939	737	2286	1875	518
5' NT of Clone Seq.	1	1	1	—			1	549	1	1
Total NT Seq.	1641	1470	604	981	1653	939	737	2286	1875	518
NT SEQ ID NO:	443	444	445	446	447	448	449	450	451	452
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	PTA- 793 09/27/99	203917 04/08/99	PTA- 181 06/07/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HCFNN75	HCF0G17	НСЕОН93	HCGBA15	HCHAC68	НСНВР49	HCHCA79	НСНС G33	HCHMY57	нсносое
Gene No.	433	434	435	436	437	438	439	440	441	442

Last AA of ORF	410	99	99	59	83	61	36	61	7	11
First I AA of A Secreted Portion C	3	48	48	19	24	27	12	17		
	2	47	47	18	23	26	11	16		
First AA of Sig Pep	1	1	1	1	-	1	1	1	1	1
AA SEQ ID NO: Y	3050	3051	3052	3053	3054	3055	3056	3057	3028	3059
	24	1321	1321	1050 3053	228	90	309	168	241	2671 3059
5' NT of of of Start AA of Codon Signal Pep	24	1321	1321			96		168	241	2671
3' NT of Clone Seq.	1393	2163	2163	1563	682	907	1508	1003	829	2841
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	25	966	966	574	-		1	-		2461
Total NT Seq.	1413	2163	2163	1588	682	907	1508	1003	879	3281
SEQ NO:	453	454	455	456	457	458	459	460	461	462
Vector	pSport1	pSport1	pSport1	Lambda ZAP II	Lambda ZAP II	Uni-ZAP XR	Uni-ZAP XR	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99	PTA- 181 06/07/99	203917 04/08/99	203917 04/08/99		203917 04/08/99	203979 04/29/99
cDNA Clone ID	нсноу52	нснов93	нснов93	HCLBK61	HCLCU75	HCMSA37	HCMSR07	HCNAI74	HCNCT01	HCNDR39
Gene No.	443	444	445	446	447	448	449	450	451	452

Last AA of ORF	16	4	31	∞	42	19	39	305	4	21
First AA of Secreted Portion (21	21		30		25	2		
		20	20		29		24			
	-	_	1	-	1	1	-	-	-	-
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	3060	3061	3062	3063	3064	3065	3066	3067	3068	3069
5' NT of First AA of Signal Pep		1011	23	268	46	385	113	140	253	395
5' NT of Start Codon	309	1011	23		46	385	113	140	253	395
3' NT of Clone Seq.	870	1725	509	917	676	1192	862	1388	769	1909
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	771	-	-	-	100	1	620	1	1
Total NT Seq.	870	1735	509	917	929	1232	862	1388	692	2629
SEQ NO:	463	464	465	466	467	468	469	470	471	472
Vector	pBluescript	pBluescript	pBluescript	pBluescript	Uni-ZAP XR	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II
ATCC Deposit No.Z and Date	203917 04/08/99	PTA- 1838 05/09/00	203917	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	
cDNA Clone ID	HCNSD91	HCNSF01	HCNSG06	HCNSG32	HCPAE41	HCQAK36	HCQAQ47	HCQAS72	нсовм95	нсосм95
Gene No.	453	454	455	456	457	458	459	460	461	462

Last AA of ORF	21	25	25	39	6	4	39	46	95	194	32
First Last AA of AA Secreted of Portion ORF				21		22	39	21	34	33	
				20		21	38	20	33	32	
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	_		_	1	1	1	1		-	-	-
AA SEQ ID NO: Y	3070	3071	3072	3073	3074	3075	3076	3077	3078	3079	5198
	481	516	516	72	292	142	175	199	394	87	41
S' NT of of Of Start AA of Codon Signal Pep	481	516	516	72	292	142	175	199	394	87	41
	1996	1414	1412	884	875	753	069	1512	2100	846	1122
5' NT 3' NT of Olone Clone Seq. Seq.	183	396	396	1	1	1	1	104	318	45	-
Total NT Seq.	2017	1414	1412	884	875	753	069	1512	2120	846	1122
SEQ SEQ NO:	473	474	475	476	477	478	479	480	481	482	2601
Vector	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203979 04/29/99	203979	203917 04/08/99	203959	203917	203918 04/08/99	203917 04/08/99	203979 04/29/99	203917	203917 04/08/99
cDNA Clone ID	нсосм95	нсосу23	нсосу23	нсорр32	нсорр61	HCQDT67	HCRAI29	HCRBI79	HCRBL20	HCRBX84	HCRBX84
Gene No.	463	464	465	466	467	468	469	470	471	472	472

		<u>-</u>	1						
Last AA of ORF	22	28	58	28	443	31	31	32	51
F A Sec Po	19	29	29	29	19	15	25	16	27
	18	28	28	28	18	14	24	15	26
First Last AA AA of of Sig Sig Pep Pep	-	1	-	1	-	-		_	-
AA SEQ D NO: Y	3080	3081	3082	3083	3084	3085	3086	3087	3088
5' NT of of of Start AA of Codon Signal Pep	98	2164	2164 3082	2164	367	309	169	405	65
5' NT of Start Codon	98	2164	2164	2164	367	309	169		65
3' NT of Clone Seq.	652	2577	2577	2577	1891	1487	262	191	218
5' NT 3' NT of Clone Seq.	1	1842	1842	1842	-	1	1	1	1
Total NT Seq.	652	2909	2918	2918	1891	1487	262	773	218
SEQ BD: NO:	483	484	485	486	487	488	489	490	491
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	ZAP Express	ZAP Express	ZAP Express	ZAP Express
ATCC Deposit No.Z and Date	PTA- 794 09/27/99	PTA- 181 06/07/99	PTA- 181 06/07/99	PTA- 181 06/07/99	PTA- 792 09/27/99	203917	203917 04/08/99	203917 04/08/99	203918 04/08/99
cDNA Clone ID	HCRMA24	HCRMR35	HCRMR35	HCRMR35	HCROC18	HCUAE53	HCUAO46	HCUAT74	HCUBA28
Gene No.	473	474	475	476	477	478	479	480	481

Last AA of ORF	37	59	39	120	45	20	4	35	33	39	29
First AA of Secreted Portion (29	22	23	19	21	20	27	32	24	28	18
	28	21	22	18	20	61	26	31	23	27	17
First Last AA AA of of Sig Sig Pep Pep	-	-	-	1	1	1	1	-	1	1	1
	3089	3090	3091	3092	3093	3094	3095	3096	3097	8608	3099
	86	27	152	337	744	229	227	200	25	<u>5</u> 9	230
5' NT of of of Start AA of Codon Signal Pep	86	27			744	229	227	200		65	230
	488	1269	858	1107	1114	371	360	205	466	1545	552
5' NT 3' NT of Clone Clone Seq.	1	1	 4	-	547	1	1	1	1	-	1
Total NT Seq.	488	1269	858	1107	1114	371	360	505	499	1545	552
SEQ BD NO:	492	493	464	495	496	497	498	499	200	501	502
Vector	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203918 04/08/99	203917 04/08/99	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99
cDNA Clone ID	HCUBC45	HCUBM41	HCUBN69	HCUBY47	HCUCV66	HCUDJ41	HCUEC55	HCUEG85	HCUES35	HCUFC77	HCUFD17
Gene No.	482	483	484	485	486	487	488	489	490	491	492

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Last AA of ORF	33	48	49	34	34	21	36	4	29	49	36
First Last AA of AA Secreted of Portion ORF	22	15	25	24	24	10	36		9	20	14
Last AA of Sig Pep	21	14	24	23	23	6	35		5	19	13
First AA of Sig Pep	1	-	1	1	-	1	-	-	-	1	-
AA SEQ ID NO: Y	3100	3101	3102	3103	3104	3105	3106	3107	3108	3109	3110
5' NT of of of Start AA of Codon Signal Pep	101	90	352	295	293	280	263	80	329	27	831
5' NT of Start Codon	101	90	352	295	293		263	80			
	354	522	573	597	594	1333	391	182	587	1630	2139
of of Clone Seq.	88	Ţ	226	250	248	1	1	1	1	1	-
Total NT Seq.	009	522	573	597	594	1333	391	182	587	1630	2139
NT SEQ BD NO:	503	504	505	909	507	508	509	510	511	512	513
Vector	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express
ATCC Deposit No.Z and Date	203917 04/08/99	203917	203917 04/08/99	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HCUFD46	HCUFE33	HCUFJ09	нсоя	HCUFQ58	HCUFX08	HCUGB76	HCUGK79	HCUGQ19	HCUGR26	HCUGR86
Gene No.	493	464	495	496	497	498	499	200	501	502	503

Last AA of ORF	10	31	84	46	10	35	28	7	4	09
First AA of Secreted Portion		23	22	40		23				25
		22	21	39		22				24
First AA of Sig Pep	1	1	-	1	1	1	1	-	1	—
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	3111	3112	3113	3114	3115	3116	3117	3118	3119	3120
F S' NT AA First of SEQ AA of First AA of NO: Signal A Pep	213	20	08	115	362	142	284	166	328	223
5' NT of Start Codon	213	50	80	115		142	284	166		223
3' NT of Clone Seq.	819	174	622	1993	1094	2439	829	2295	1055	1179
5' NT 3' NT of of Total Clone Clone NT Seq. Seq. Seq.	-	1	1	40	7	1	-	-	150	∞
	819	174	622	1993	1094	2439	829	2295	1055	1179
SEQ N NO: O	514	515	516	517	518	519	520	521	522	523
Vector	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express
ATCC Deposit No.Z and Date	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203979 04/29/99	203917 04/08/99	PTA- 1838 05/09/00
cDNA Clone ID	нсине27	HCUHL82	нсинм71	HCWAK88	HCWAL10	HCWAT71	нсwвQ52	НСМСН16	нсмрм69	HCWEB38
Gene No.	504	505	909	507	208	509	510	511	512	513

Last AA of ORF	27	119	9	36	49	49	43	12	32	43
First AA of Secreted Portion	24	25		∞	19	19	28		23	19
	23	24		7	18	18	27		22	18
First AA of Sig Pep	1	-	-	1	-	-	1	-	-	-
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	3121	3122	3123	3124	3125	3126	3127	3128	3129	3130
S' NT AA of SEQ First ID AA of NO: Signal Y Pep	98	25	135	1156	227	227	149	262	190	527
5' NT of of of Start AA of Codon Signal Pep	98	25	135		227	227	149	262	190	527
3' NT of Clone Seq.	883	356	657	1901	358	358	303	810	1256	657
S' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1		1	1	12	12	T	1	1	1
	883	383	657	1901	375	375	303	810	1256	657
SEQ BD NO:	524	525	526	527	528	529	530	531	532	533
Vector	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express	ZAP Express
ATCC Deposit No.Z and Date	203917 04/08/99	PTA- 1838 05/09/00	203917 04/08/99	203917	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HCWEB72	HCWEF04	HCWEI82	нсмем96	HCWFJ16	HCWFJ16	HCWFK03	нсмнD30	HCWHT34	HCWHT52
Gene No.	514	515	516	517	518	519	520	521	522	523

A F	_ [33	99	39	30	38	243	33	44	117
Last AA of ORF	41	33	و	Ω.	Ç.	3	- 5	3	4	
First AA of Secreted Portion	24	19	46	25	19	30	15	21	18	18
Last AA of Sig Pep	23	18	45	24	18	29	14	20	17	17
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	-		1	1	1		1	1	-
AA SEQ ID NO: Y	3131	3132	3133	3134	3135	3136	3137	3138	3139	3140
5' NT of First AA of Signal Pep	146	244	609	<i>2</i> 7	173	54	419	95	178	69
5' NT of of of Start AA of Codon Signal Pep	146	244	609	27	173	54		96		69
	979	342	792	1144	2092	643	1896	3362	842	419
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.		1	998		1	-	_	-		
	979	342	808	1300	2092	643	1896	3362	842	419
NT SEQ ID NO:	534	535	536	537	538	539	540	541	542	543
Vector	ZAP Express	ZAP Express	ZAP Express	ZAP Express	pBluescript SK- 538	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917	203979 04/29/99	203917 04/08/99	PTA- 181 06/07/99	203917 04/08/99	203917 04/08/99	PTA- 794 09/27/99
cDNA Clone ID	HCWK032	HCWLE50	HCWUF93	HCWUW24	HCYBA32	HDAAV67	HDABR74	HDABW45	HDACJ52	НДСВМ09
Gene No.	524	525	526	527	528	529	530	531	532	533

Last AA of ORF	39	32	21	20	85	<i>L</i> 9	28	39	54	38	179
First I AA of Secreted Portion C	18	21	17	32	19	22	21	16	35	21	16
	17	20	70	31	18	21	20	15	34	20	15
First Last AA AA of of Sig Sig Pep	-	-	_	1	-	-	1	1		~	-
AA ISEQ ID NO:	3141	3142	3143	3144	3145	3146	3147	3148	3149	3150	3151
	155		53	93	414	294	302	249	146	210	192
5' NT of of of Start AA of Codon Signal Pep	155	357	53	93	414	294	302	249	146	210	192
	1262	1624	224	1567	1681	<i>L</i> 69	733	089	1661	1336	3569
5' NT 3' NT of of Clone Clone Seq. Seq.	1	1	-		1	-	-	1	1	71	548
Total NT Seq.	1262	1624	224	1567	1681	269	733	089	1661	1336	3569
SEQ NO.	544	545	546	547	548	549	550	551	552	553	554
Vector	pCMVSport 2.0										
ATCC Deposit No.Z and Date	203917		203917	203917	203979	203917	203917	203917	203917	203917	203979 04/29/99
cDNA Clone ID	HDFAB86	HDFIB37	HDFMB91	HDHAA55	HDHEA33	HDHEB12	HDHEB80	HDHIA16	HDHIA26	HDHMA71	HDLAL94
Gene No.	534	535	536	537	538	539	540	541	542	543	544

Last AA of ORF	81	55	59	31	33	24	43	32	26	63
if Li	<u>~</u>	3		-						
First AA of Secreted Portion	23	25		21	20	23		32	22	44
Last AA of Sig Pep	22	24		20	19	22		31	21	43
First Last AA AA of of Sig Sig Pep	1	1	1	1	-	1		—	_	-
AA SEQ ID NO: Y	3152	3153	3154	3155	3156	3157	3158	3159	3160	3161
	389	181	06	371	126	240	95	305	320	170
5' NT of of of Start AA of Codon Signal Pep	389	181	06	371	126	240	95	305	320	170
	2074	2010	1426	2379	1735	1581	1226	3840	2243	1635
5' NT 3' NT of Olone Clone Seq. Seq.			-	247	1	1	1		-	11
Total NT Seq.	2074	2010	1426	2382	1735	1581	1226	3840	2243	1635
SEQ NO:	555	556	557	558	559	560	561	562	563	564
Vector	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	CMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0 561	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0
ATCC Deposit No.Z and Date	PTA- 1838 05/09/00	203917 04/08/99	203917 04/08/99	203917 04/08/99	203960 I	203960 04/26/99	203960 104/26/99	203960 04/26/99	PTA- 791 09/27/99	203960 04/26/99
cDNA Clone ID	HDPAB86	HDPAE80	HDPAQ86	HDPBD56	HDPBN48	HDPCG79	HDPCV29	HDPDA36	HDPDC59	HDPFG13
Gene No.	545	546	547	548	549	550	551	552	553	554

Last AA of ORF	34	42	100	41	8	30	09	38	21	57
First AA of Secreted Portion (15	91	20	7		20	36	20	∞	37
	14	15	16	9		19	35	19	7	36
First Last AA AA of of Sig Sig Pep Pep	1	1	1	1	-	-			_	П
AA SEQ ID NO: Y	3162	3163	3164	3165	3166	3167	3168	3169	3170	3171
····	283	172	17	1690	1202 3166	96	169	212	102	38
5' NT of of of Start AA of Codon Signal Pep		172	17		1202	06	169	212		38
	1533	2185	1119	1756	2321	1371	4115	2251	826	1646
5' NT of Clone Seq.	1	П	П	29	868	_	-		-	1
Total NT Seq.	1533	2185	1119	2608	2322	1371	4115	2251	1011	1646
SEQ NO:	595	995	267	268	695	570	571	572	573	574
Vector	pCMVSport 3.0	pCMVSport 3.0 571	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0 574					
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	203979 04/29/99	203918 04/08/99	203960 04/26/99	PTA- 794 09/27/99	PTA- 793 09/27/99	203959 04/26/99	203979 04/29/99
cDNA Clone ID	HDPFK27	HDPFZ05	HDPGA84	HDPGR80	HDPGU14	HDPGX09	HDPIE44	HDPIE73	HDPIF35	HDPIF65
Gene No.	555	556	557	558	559	260	561	562	563	564

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Last AA of ORF	51	35	186	57	23	453	21	78	32	35
First AA of Secreted Portion	46	17	14	17		29		51	25	16
	48	16	13	16		28		20	24	15
AA First Last SEQ AA AA Of Of Of Sig Sig Y Pep Pep			-	1	1	1	-	1	-	1
AA SEQ NO:	3172	3173	3174	3175	3176	3177	3178	3179	3180	3181
5' NT of First AA of Signal		268	603	145	382	102	274	214	86	332
of of Start Codon	181	268		145		102	274	214	86	332
	2729	1978	1348	2816	1250	1795	2486	554	1422	1181
5' NT of Clone Seq.		1	12	1	1			-	-	
Total NT Seq.	2729	1978	1990	2816	1250	1795	2486	554	1422	1181
SEQ NO:	575	576	577	578	579	580	581	582	583	584
Vector	pCMVSport 3.0	pCMVSport 3.0 576	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0
ATCC Deposit No.Z and Date	203960	PTA- 793			203960 04/26/99	203960 04/26/99	PTA- 794 09/27/99	203979 04/29/99	203960	203960 04/26/99
cDNA Clone ID	HDPIH25	HDPIY31	HDPJH72	HDPJV53	HDPJV75	HDPKC55	HDPKD16	HDPMC52	HDPML04	HDPMM22
Gene No.	595	995	292	995	995	270	571	572	573	574

Last AA of ORF	89	20	∞	97	310	49	16	16	31
First Last AA of AA Secreted of Portion ORF		21		18	21	50		41	25
	6	20		17	20	19		13	24
First AA of Sig Pep	-	-	-	-		-	1		-
	3182	3183	3184	3185	3186	3187	3188	3189	3190
5' NT of First AA of Signal Pep	40	422	446	691	257	271	303	252	186
5' NT of of of Start AA of Codon Signal Pep		422	446		257	271	303	252	186
	1430	1719	797	1868	2414	686	1112	1254	1240
s' NT of Clone Seq.	1	394		551	82	1	1	-	
Total NT Seq.	1430	1719	797	1868	2444	989	1112	1254	1240
SEQ NO:	585	586	587	588	589	590	591	592	593
Vector	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0
ATCC Deposit No.Z and Date	203960 04/26/99	PTA- 181 06/07/99	PTA- 794 09/27/99	PTA- 1838 05/09/00	PTA- 791 09/27/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203979 04/29/99
cDNA Clone ID	HDPNC21	HDPNJ26	HDPOD73	HDPOT33	HDPPB70	HDPPC19	HDPPE05	HDPSA70	HDPSS56
Gene No.	575	576	577	578	579	580	581	582	583

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Last AA of ORF	39	39	39	35	37	23	310	310	370	137	0
First AA of Secreted Portion	28	28	28	18	27	23	20	50	20	32	
	27	27	27	17	26	22	19	19	19	31	
First AA of Sig Pep	1	-	1	1	1	1	1	-	-	-1	
	3191	3192	3193	3194	3195	3196	3197	3198	3199	3200	3201
	244	244	244	182	236	301	469	469	470	96	728
5' NT of of of Start AA of Codon Signal Pep	244	244	244	182	236	301	469	469	470	96	728
	1009	1009	1009	1724	1519	1108	1437	1437	1439	1960	1077
S' NT 3' NT of of Clone Clone Seq. Seq.		1	1	-	1	33	53	53	53	-	482
Total NT Seq.	1337	1337	1337	1724	1519	1108	1579	1561	1581	1960	1077
NT SEQ D NO:	594	595	596	597	598	599	009	601	602	603	604
Vector	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0 600	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0 604
ATCC Deposit No.Z and Date	203960	203960 04/26/99	203960	203979	203960	203960	203960	203960		203960	203960 04/26/99
cDNA Clone ID	HDPSZ07	HDPSZ07	HDPSZ07	HDPTI49	HDPTP22	HDPYE25	нросрое	нросрое	нросрое	HDQGN08	HDQG062
Gene No.	584	585	586	587	588	589	290	591	592	593	594

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Last AA of ORF	39	50	20	32	55	23	49	29	126	52
First AA of Secreted Portion	27	28	23	25	18	21	22	12	29	22
Last AA of Sig Pep	26	27	22	24	17	20	21	11	28	21
First AA of Sig Pep	—	1	1	-	1	1	-	-	-	-
AA SEQ ID NO: Y	3202	3203	3204	3205	3206	3207	3208	3209	3210	3211
5' NT AA of SEQ First ID AA of NO: Signal Y	473	77	65	11	239	226	208	351	926	199
5' NT of of of Start AA of Codon Signal Pep	473	77	65	11	239	226			926	199
	2054	788	782	1387	545	924	1433	477	1417	1267
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-	1	1	1	1		_	1	851	1
Total NT Seq.	2054	788	782	1387	545	924	1433	477	1857	1267
SEQ SEQ NO:	909	909	209	809	609	610	611	612	613	614
Vector	pCMVSport 3.0	pCMVSport 2.0 606	pCMVSport 2.0	Uni-ZAP XR	pCMVSport 2.0					
ATCC Deposit No.Z and Date	PTA- 793	203960 04/26/99	203960 04/26/99	203960	203960	203960	203979	203960	203960	203979 04/29/99
cDNA Clone ID	HDQPM16	HDRAA17	HDRAC68	HDSAC78	HDSAH37	HDSAM57	HDSA014	HDSA064	HDSAP15	HDTAR39
Gene No.	595	296	597	298	599	009	601	602	603	604

Last AA of ORF	20	37	34	37	75	35	25	6	37	36
First AA of Secreted Portion	47	31	31	21	34	14			34	16
Last AA of Sig Pep	46	30	30	20	33	13			33	15
AA First SEQ AA ID of NO: Sig Y Pep	1	-	—	-	-	1	1	1	_	-
AA First SEQ AA ID of NO: Sig Y Pep	3212	3213	3214	3215	3216	3217	3218	3219	3220	3221
5' NT of First AA of Signal Pep	62	203	348	105	223	410	443	19	156	250
5' NT of of of Start AA of Codon Signal Pep	62	203	348	105	223	410		16	156	250
3' NT of Clone Seq.	915	1358	1335	932	<i>L</i> 69	611	9/9	572	2235	1476
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	1	1	1	1	1	-	-
	915	1358	1335	932	269	611	9/9	572	2235	1476
SEQ NO:	615	919	617	618	619	620	621	622	623	624
Vector	pCMVSport 2.0 615	pCMVSport 2.0 616	pCMVSport 2.0	pCMVSport 2.0	203960 pCMVSport 2.0 04/26/99	203960 pCMVSport 2.0 620 04/26/99	pCMVSport 2.0	pCMVSport 2.0	pCMVSport 2.0	pCMVSport 2.0 624
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99	
cDNA	HDTAS57	HDTBP62	HDTBQ77	HDTDA48	HDTDE66	HDTDG75	HDTDS09	HDTFF53	HDTGW76	HDTGZ56
Gene No.	909	909	209	809	609	610	611	612	613	614

Last AA of ORF	31	85	207	33	34	31	31	33	55
First AA of Secreted Portion		30	23	13	32	27	27	16	26
Last AA of Sig Pep		29	22	12	31	26	26	15	25
First Last AA AA of of Sig Sig Pep Pep	1		1	1	-	_	1	-	1
AA SEQ ID NO: Y	3222	3223	3224	3225	3226	3227	3228	3229	3230
5' NT AA of SEQ of First ID Start AA of NO: Codon Signal Y	114	401	112	294	35	1869 3227	3022 3228	14	144
	114	401	112	294	35	1869	3022	14	
3' NT of Clone Seq.	596	1735	1388	887	804	3206	4359	324	765
5' NT of Clone Seq.	1	1	1	1	1	3264 1713	2866	-	1
Total NT Seq.	969	1735	1388	887	804	3264	4417	324	765
NT SEQ ID NO:	625	979	627	628	679	630	631	632	633
Vector	pCMVSport 2.0	pCMVSport 2.0	pCMVSport 2.0	pSport1	pSport1	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 793 09/27/99	PTA- 793 09/27/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99	PTA- 181 06/07/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	S8ZHLQH	HDTM39	HDTKJ29	HDUAB12	HDUAD68	HE2AC74	HE2AC74	HE2AC75	HE2AI94
Gene No.	615	616	617	618	619	620	621	622	623

First Last AA AA First L of of AA of A Sig Sig Secreted Pep Pep Portion	1 16 17 20	3 1 13 14 59	4 1 14 15 30			5 1 28 29 34	6 1 20 21 44	2	7 1 14 15 16		18 1 28 29 68	1 27 28 38		50 1 29		51 1 10	
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	95 3242	842 3243	175 3244			493 3245	101 2776	101	213 3247		117 3248		90 3249	374 3250		79 3251	
7	1521 95	1185	1640 175			2191 493	\perp	1593 181	014 213		1609 117		526 96	1582 374		968 79	
5' NT of Clone Seq.		563	-	-		279		-	† -	-	-	•	526 77	1500	1 700	1 890	1 00
NT SEQ ID Total NO: NT X Seq.	645 1521	646 1185		647 1049		648 3484		649 1593	-	656 069	\neg	0.01	652	(2)	600	1733	0.04
Vector	Uni-ZAP XR	IIni-7AP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	1	Uni-ZAP XR	- 1	Uni-ZAP AK	Uni-ZAP XR	- 1	Uni-ZAP XK	_	Uni-ZAP AK
ATCC Deposit No.Z and Date	203960	04/26/99	04/26/99	PTA-	791	203918	04/08/99	203979	04/29/99	203960	04/26/99	203918	203960	04/26/99	203960	04/26/99	203960
cDNA Clone ID		- 1	HE2DJ84	HE2DY23		THOUNDS	HEZD 123	HE2EE80		HE2EH45		HE2FE89	HE2FR49		HE2GB19		HE2G081
Gene No.	635	3	929	637		95,	038	639	<u> </u>	640		641	642	! 	643		644

Last AA of ORF	31	36	7.5	23	14	118	35	-	23	113	42
First I AA of Secreted Portion (29		19	14		30	10		18	15	26
	78	!	18	13		29	6		17	14	25
First AA of Sig Pep	-	-		1	1	1	1	_	-		-
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	3252	3253	3254	3255	3256	3257	3258	3259	3260	3261	3262
	. 1		111	1331 3255	861	101	l → l	79	19	69	142
5' NT of of of Start AA of Codon Signal Pep	154	225	111		861	101		79	19	69	142
	755	1875	1222	2048	1746	516	1671	1356	088	1003	1061
5' NT 3' NT of Olone Clone Seq.	1	1		1	601	1	1	1	-	-	135
Total NT Seq.	755	1875	1222	2048	1746	516	1671	1356	880	1003	1061
SEQUENCE NO SEQUEN	655	959	129	658	629	099	199	662	663	664	999
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960	203960 04/26/99	203960	203960	203918	203960	203960 04/26/99	203960	203979	203979	203960
cDNA Clone ID	HE2HB61	HE2HB64	HE2HF76	HE2ID09	HE2IE66	HE2NW57	HE20A95	HE2OC39	HE2PB61	HE2PI43	HE2PJ56
Gene No.	645	646	647	648	649	059	651	652	653	654	655

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Last AA of ORF	10	23	13	98	35	17	25	18	118	14	380
First AA of Secreted Portion		16		21	26	15	16		17	11	21
Last AA of Sig Pep		15		20	25	14	15		16	10	20
First AA of Sig Pep	—	1	1	1	1	1	1	П	—	-	-
AA SEQ ID NO: Y	3263	3264	3265	3266	3267	3268	3269	3270	3271	3272	3273
	231	107	257	48	9/1	364	285	89	289	202	208
5' NT of of of Start AA of Codon Signal Pep		107	257	48	9/1		285	89	289		208
	1080	464	1708	603	1415	774	3334	918	2270	1859	2867
5' NT 3' NT of of Clone Clone Seq. Seq.	1	1	1	-	1	8	1	1	1	_	1
Total NT Seq.	1080	464	1708	603	1415	780	3334	918	3193	1859	2867
SEQ BD NO:	999	<i>L</i> 99	899	699	0/9	671	672	673	674	675	9/9
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	203979	203960 04/26/99	203960 04/26/99	203918 04/08/99	203960	203960 04/26/99	203918 04/08/99	203979 04/29/99
cDNA Clone ID	HE6CJ41	HE6DC37	HE6DN83	HE6EI30	HE6ET70	HE6G065	HE8AN83	HE8AU68	HE8BE20	HE8BP05	HE8BP64
Gene No.	959	657	658	629	099	661	662	699	664	999	999

Last AA of ORF	11	31	28	105	22	39	28	36	22	32
First AA of Secreted Portion		19		27		18		16		18
Last AA of Sig Pep		18		26		17		15		17
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep	1	-	<u></u>	1	-		<u> </u>	-	1	1
AA SEQ ID NO: Y	3274	3275	3276	3277	3278	3279	3280	3281	3282	3283
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	133	207	168	2371	244	111	163	347	235	137
5' NT of Start Codon	133	207	168	2371		111	163	347	235	137
	1875	1651	2532	3533	1902	1322	2148	2556	1642	1783
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	12	-	-	2277		-	-			1
_	1875	1651	2292	3560	1902	1538	2148	2608	1642	1783
NT SEQ ID NO:	<i>LL</i> 9	8/9	629	089	681	682	683	684	985	989
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR							
N _O	Uni-2	Uni-2	Uni-2							
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99							
cDNA Clone ID	HE8BQ49	HE8BR18	HE8BR30	HE8BT58	HE8BU60	HE8CA13	HE8CC34	НЕ8СН08	HE8DG02	HE8DK52
Gene No.	<i>L</i> 99	899	699	029	671	672	673	674	5/9	929

Last AA of ORF	38	21	71	145	32	1117	30	20	14	37
First AA of Secreted Portion	28	15	17	27		24	27	70		23
Last AA of Sig Pep	27	14	16	56		23	26	19		22
First AA of Sig Pep		П	-	-	-	-	1			1
AA SEQ ID NO: Y	3284	3285	3286	3287	3288	3289	3290	3291	3292	3293
5' NT of First AA of Signal Pep	29	65	914	315	138	172	187	251	223	1206 3293
5' NT of Start Codon	29	99	914		138	172	187	251	223	1206
	1799	3198	2576	1054	2472	1606	2505	1271	1748	3685
5' NT 3' NT of of Clone Clone Seq. Seq.		П	39	П	2	83	28	П	-	1171
Total NT Seq.	1799	3198	4185	1054	2472	1606	2505	1271	1748	3707
SEQ BD NO:	<i>L</i> 89	889	689	069	691	769	693	694	569	969
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 791 09/27/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99	203960 04/26/99	203918 04/08/99	203979 04/29/99
cDNA Clone ID	HE8DZ94	HE8EN79	HE8EX86	HE8FC10	HE8FG15	HE8FG24	HE8FK78	HE8FL24	HE8FL68	HE8FR53
Gene No.	<i>LL</i> 9	829	629	089	681	682	683	684	985	989

	1		10.77		1					
Last AA of ORF	99	8	37	9	32	13	282	35	62	28
First Last AA of AA Secreted of Portion ORF	45	23	26		24		25		25	21
	44	22	25		23		24		24	20
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1		-	1	1	1	-	1		1
9 1 —	3294	3295	3296	3297	3298	3299	5199	3300	3301	3302
5' NT of of of Start AA of Codon Signal Pep	06	276	127	1387	61	74	419	837	4	47
5' NT of Start Codon	90	276	127	1387	61	74	419	837	4	47
3' NT of Clone Seq.	1307	2303	2719	2600	2721	2626	3357	997	1589	3161
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.		12	1		-			574	1	1
Total NT Seq.	1307	2304	2719	2600	2721	2626	3357	1034	1589	3161
SEQ BD NO:	<i>L</i> 69	869	669	700	701	702	2602	703	704	705
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	PTA- 793 09/27/99	203960 04/26/99	203979 04/29/99	PTA- 181 06/07/99	PTA- 181 06/07/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HE8MA27	HE8MG56	НЕ8МQ01	HE8MS43	HE8MY77	HE8NC81	HE8NC81	HE8NO09	HE8QU21	HE8SH74
Gene No.	289	889	689	069	169	692	692	693	694	695

Last AA of ORF	74	18	30	26	20	34	16	27	89	12
First AA of Secreted Portion	23	17	18	19		23	15		11	
Last AA of Sig Pep	22	16	17	18		22	14		16	
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	1	-
AA SEQ ID NO: Y	3303	3304	3305	3306	3307	3308	3309	3310	3311	3312
5' NT of SEQ by Start AA of NO: Start AA of NO: Codon Signal Y F	142	654	188	1629	\$8	9	139	152	295	701
5' NT of Start Codon	142	654	188		\$8	9	139	152	295	701
3' NT of Clone Seq.	1409	1931	1934	2283	2742	1294	930	1393	1913	2443
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	8	476	-	1696	1		1	1	_	400
Total NT Seq.	1409	1931	2128	2283	2742	1294	930	1393	1913	2502
NT SEQ ID NO:	902	707	708	709	710	711	712	713	714	715
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	PTA- 181 06/07/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960	203960 04/26/99	203979 04/29/99	203960	PTA- 793 09/27/99
cDNA Clone ID	HE8UX34	HE9AE05	HE9BJ14	HE9CI81	HE9CJ38	HE9CM11	HE9CN58	HE9CV59	HE9DG54	нЕ9DН59
Gene No.	969	<i>L</i> 69	869	669	700	701	702	703	704	705

			r	r						
Last AA of ORF	44	9	26	30	ς.	5	9	31	109	19
First AA of Secreted Portion	36		25	21				21	27	
ast AA of Sig	35		24	20				20	26	
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	-	1	-	1	1	1	1
AA SEQ ID NO: Y	3313	3314	3315	3316	3317	3318	3319	3320	3321	3322
5' NT of First AA of Signal Pep	239	541	125	159	1949	739	412	200	09	276
5' NT of Start Codon	239		125	159	1949	739	412	200	09	276
5' NT 3' NT of of of Total Clone Clone NT Seq. Seq.	1219	1279	1086	1276	2735	1525	1614	937	1329	2455
5' NT 3' NT of Ol Clone Clone Seq.	1	318	1	1	1804	594		1	10	1
Total NT Seq.	1276	1279	1086	1276	2757	1547	1614	937	1329	2455
NT SEQ ID T NO:	716	717	718	719	720	721	722	723	724	725
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99						
cDNA Clone ID	HE9DZ47	НЕ9ЕС36	HE9EM54	НЕ9FH28	HE9HE13	не9не13	HE9HF59	HE9HV71	HE9NB82	HE9NE43
Gene No.	902	707	708	709	710	711	712	713	714	715

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Last AA of ORF	26	36	63	5	30	10	14	41	15	15
First AA of Secreted Portion	23	29	16					19		13
	22	28	15					18		12
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	-	-	_	-	-	-	-	-	-	-
AA SEQ ID NO: Y	3323	3324	3325	3326	3327	3328	3329	3330	3331	3332
5' NT of First AA of Signal Pep	515	2540 3324	89	299	178	107	296	219	217	377
5' NT of of of Start AA of Codon Signal Pep	515	2540	89		178	107	296			377
3' NT of Clone Seq.	834	3359	1094	1243	818	847	662	2254	1079	2166
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	2187	-	П		-	-		_	-
	834	3371	1094	1243	818	847	662	2254	1079	2166
NT SEQ ID NO:	726	727	728	729	730	731	732	733	734	735
Vector	Uni-ZAP XR									
ATCC Deposit No.Z and Date	203960 04/26/99	PTA- 181 06/07/99								
cDNA Clone ID	HE9RN58	HE9TA42	HEAAC21	HEAAC39	HEAAC48	HEAAD63	HEAAE19	HEAAM54	HEAAM96	HEAAN52
Gene No.	716	717	718	719	720	721	722	723	724	725

Last AA of ORF	40	53		37	55	19	59	20	62	40
	4	5		<u></u>	5		2	ζ.	9	4
First AA of Secreted Portion	36	24		35	25		11	15	28	20
Last AA of Sig Pep	35	23		34	24		10	14	27	19
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep	1	-	1	1	-	1	1			1
AA] SEQ ID NO: Y	3333	3334	3335	3336	3337	3338	3339	3340	3341	3342
5' NT AA 5' NT of SEQ of First ID Start AA of NO: Codon Signal Y Pep	73	196	189	1060 3336	216	590	280	322	1080 3341	1461
5' NT of Start Codon	73	196	189		216				1080	1461
	632	1104	924	1492	638	933	408	1687	1266	2710
5' NT of Clone Seq.	-		-	898	П	298	-	-	30	1273
Total NT Seq.	632	1104	924	1492	638	944	408	1687	1266	2902
NT SEQ D NO:	736	737	738	739	740	741	742	743	744	745
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203918 04/08/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99	203960 04/26/99
cDNA Clone ID	HEAAU28	HEAAW54	HEAAW94	HEBAP51	HEBAT05	HEBBF78	HEBBK04	HEBCN80	HEBCW57	HEBDF90
Gene No.	726	727	728	729	730	731	732	733	734	735

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Last AA of ORF	142	43	36	81	7	51	40	17	41	7
First Last AA of AA Secreted of Portion ORF	22	26	22	32		20	33		17	
Last AA of Sig Pep	21	25	21	31		19	32		16	
AA First SEQ AA ID of NO: Sig Y Pep		_	1	1	1	1	1	1	1	1
AA SEQ NO:	3343	3344	3345	3346	3347	3348	3349	3350	3351	3352
5' NT of First AA of Signal	Pep 734	381	201	153	615	158	26	1157	38	291
5' NT of Start Codon	734	381	201	153	615	158	26	1157	38	291
3' NT of Clone Seq.	1305	590	752	419	949	440	1504	1607	2141	2414
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	486	-	1	_	380	_	6	942	-	1
Total NT Seq.	1328	590	752	419	949	440	1504	1635	2141	2414
SEQ NO:	746	747	748	749	750	751	752	753	754	755
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960	203979 04/29/99	203918 04/08/99	203960 04/26/99	PTA- 792 09/27/99	203960 04/26/99
cDNA Clone ID	HEBDW31	HEBFL36	HEBGC01	HEBGE23	HEBGE85	HEBGJ94	HEBGM06	HEEAB58	HEEAF49	HEEAJ46
Gene No.	736	737	738	739	740	741	742	743	744	745

Last AA of ORF	57	52	38	22	15	10	43	89	26	10	218
First I AA of A Portion C	35	19	36	22			37	16			21
	34	18	35	21			36	15		:	20
First AA of Sig Pep		1	1	1	1	1	1	1	1	1	1
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	3353	3354	3355	3356	3357	3358	3359	3360	3361	3362	3363
5' NT of First AA of Signal Pen		992	188	215	182	37	1121	316	712	75	165
5' NT of of of Start AA of Codon Signal Pen	1111	992	188	215		37	1121		712	75	165
	926	1336	626	2105	1491	1460	2448	968	2070	695	1008
S' NT 3' NT of of of Clone Clone NT Seq. Seq. Seq.	1	089	-		1	1	296	1	49	-	119
Total NT Seq.	929	3940	626	2105	1491	1460	2653	968	2070	569	1123
SEQ NO: DE	756	757	758	759	760	761	762	763	764	292	992
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203959 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HEGAI20	HEIAC52	HELAC55	HELAT58	HELAW94	HELDF80	негрн39	HELDK79	HELDQ42	HELEE85	HELEL76
Gene No.	746	747	748	749	750	751	752	753	754	755	756

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Last AA of ORF	195	21	18	41	21	29	45	38	38	30
First AA of Secreted Portion	21		7	17	13	21	37	22	22	16
Last AA of Sig Pep	20		9	16	12	20	36	21	21	15
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep	1	1	1	1	-	-		1	-	1
AA SEQ ID NO: Y	3364	3365	3366	3367	3368	3369	3370	3371	3372	3373
5' NT of First AA of Signal Pep	42	08	118	209	255	171	935	293	293	169
5' NT of of Of Start AA of Codon Signal Pep	42	80		209	255	171	935	293	293	169
	1255	1965	1901	2354	2298	1296	3147	1302	1483	1443
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	292	-	-	842	T	П	1
	1255	1965	1901	2354	2298	1296	3147	1432	1483	1443
NT SEQ ID NO:	191	89/	692	170	771	772	773	774	775	9//
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR						
ATCC Deposit No.Z and Date	203960 04/26/99	203979 04/29/99	203959 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 793 09/27/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99
cDNA Clone ID	9/1313Н	HELEO45	HELFA57	нег гозо	HELGF28	HELGP60	HELHN47	неснри	HELHP11	HEMAE30
Gene No.	757	758	759	760	761	762	763	764	765	992

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Last AA of ORF	35	78	∞	34	92	12	49	20	30	34	28
First AA of Secreted Portion	28	25		26	21		32		17	26	22
Last AA of Sig Pep	27	24		25	70		31		16	25	21
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	-	-	1			-	1		1	1
AA SEQ ID NO:	3374	3375	3376	3377	3378	3379	3380	3381	3382	3383	3384
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	276	323	270	66	92	234	21	148	216	84	167
5' NT of Start Codon	276	323	270	66	92	234	21	148	216	84	
3' NT of Clone Seq.	1213	2512	1356	850	1018	1693	1136	2405	2937	1709	1885
S' NT 3' NT of of Clone Clone NT Seq. Seq.	1	195	П	П	-	-	-	_	-	-	1
Total NT Seq.	1213	2667	1356	850	1018	1693	1136	2405	2937	1709	1885
SEQ BO: NO:	LLL	778	622	780	781	782	783	784	785	982	787
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203979 04/29/99	203960	203960	203960 04/26/99	203960 04/26/99	203979 04/29/99	203960	203960	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HEMBV40	HEMCJ80	HEMCL55	HEMDB07	HEMDR05	HEMGK71	HEOMF59	НЕОМ173	HEOMR67	HEOMU25	HEOMU44
Gene No.	191	292	692	770	771	772	773	774	775	9//	777

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Last AA il of ORF	37	73	41	∞	40	37	91	48	43
First AA of Secreted Portion	26	14	17	7	18	23	26	25	26
Last AA of Sig Pep	25	13	16	9	17	22	25	24	25
AA First SEQ AA ID of NO: Sig Y Pep			-	-	-	-	-	-	1
AA SEQ ID NO: Y	3385	3387	3388	3389	3390	3391	3392	3393	3394
5' NT AA of SEQ First ID AA of NO: Signal Y Pep	136		223	123	294	259	1218	22	843
5' NT of Start Codon	136	38	223		294	259	1218	22	
3' NT of Clone Seq.	1078	1258	346	269	464	453	2212	1518	1498
5' NT 3' NT of of Clone Clone Seq. Seq.	1	1	1	220	-	-	1185	П	1
Total NT Seq.	1078	1258	346	541	464	453	2212	1518	1498
NT SEQ D NO:	788	790	791	792	793	794	795		
Vector	pSport1		Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 3.0 796	pCMVSport 3.0 797
ATCC Deposit No.Z and Date	203960 04/26/99 203960	04/26/99 203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 791 09/27/99	203960 04/26/99
cDNA Clone ID	HEONI85	1	HEPAD15	HEPBC23	HEPBV09	HEPCF35	HEPCU48	НЕОАН47	НЕQАР92
Gene No.	778	780	781	782	783	784	785	786	787

Last AA of ORF	38	19	19	19	134	35	24	39	12	13
First I AA of AS of Portion C	33				51	23	18	19		
	32				50	22	17	18		
First AA of Sig Pep	-	_	-	-	1	1		1	1	-
AA 1 SEQ ID NO: Y	3395	3396	3397	3398	3399	3400	3401	3402	3403	3404
5' NT AA of SEQ First ID AA of NO: Signal Y	130	505	2603	2603 3398	92	81	227	95	86	453
5' NT of of of Start AA of Codon Signal Pep	130	505	2603	2603	92	81	227	95		453
3' NT of Clone Seq.	1016	662	2731		1025	920	815	286	246	1701
5' NT 3' NT of of Of Clone Clone NT Seq. Seq.	15	1	2346	2791 2346 2731	1	1	1	1	П	_
Total NT Seq.	1626	699	2791	2791	1025	920	815	985	246	1701
SEQ NO:	798	662	008		802	803	804	805	908	807
Vector	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	203960 pCMVSport 3.0 801 04/26/99	pCMVSport 3.0	pCMVSport 3.0	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99		203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99
cDNA Clone ID	HEQAV53	HEQBJ01	HEQBJ01	HEQBJ01	НЕОВМ94	неосв93	HERAI63	HERAQ22	HERAS61	HESAG57
Gene No.	788	789	790	791	792	793	794	795	962	797

Last AA of ORF	43	17	74	435	32	32	27	14	105	36
First AA of Secreted Portion	26		42	22	25	25			16	25
Last AA of Sig Pep	25		41	21	24	24			15	24
First Last AA AA of of Sig Sig Pep	1	1	_	1	1	1	1		1	-
AA SEQ ID NO: Y	3405	3406	3407	3408	3409	3410	3411	3412	3413	3414
5' NT of First AA of Signal Pep	1198	263	183	14	148	148	64	177	1323	195
5' NT of of of Start AA of Codon Signal Pep	1198		183	14	148	148	64	177	1323	195
3' NT of Clone Seq.	2458	728	1697	2047	1805	1804	1238	2272	2810	1755
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1019	14	1	-	2	2	1	56	1117	64
Total NT Seq.	2593	728	1697	2047	1805	1804	1238	2272	2811	1758
SEQ BD NO:	808	608	810	811	812	813	814	815	816	817
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99	PTA- 181 06/07/99	203979 04/29/99	203979 04/29/99	203960	203979 04/29/99	203960 04/26/99	203979 04/29/99
cDNA Clone ID	HETAA62	HETBB70	HETBJ88	HETCM67	HETDD61	HETDD61	HETDJ34	HETDM73	HETDP76	HETFO57
Gene No.	262	799	008	801	802	803	804	805	908	807

Last AA of ORF	21 29	30	51	30	30	9	39	56	4
First AA of Secreted Portion (21	17	20	19		23	27	26
		20	16	61	18		22	56	25
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep		1		1	-	1	1	1	1
AA SEQ ID NO: Y	3415 3416	3417	3418	3419	3420	3421	3422	3423	3424
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	127	121	252	16	344	233	236	52	153
5' NT of Start Codon	229	121		16	344	233	236	52	153
3' NT of Clone Seq.	1918	096	615	1095	513	962	2238	499	788
S' NT 3' NT of of Of Clone Clone NT Seq. Seq.	1 172		1	-	182	1	1	-	1
	1918 1817	096	636	1095	513	962	2238	499	788
SEQ BD NO:	818 819	820	821	822	823	824	825	826	827
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203979 04/29/99 203960	203960	203960 04/26/99	203960 04/26/99	203960	203960 04/26/99	PTA- 1838 05/09/00	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HETGZ31 HETHD26	HETHM27	HETIN36	HFAAI17	HFAAJ45	HFADF41	HFADM09	HFAUA23	HFCAG75
Gene No.	808	810	811	812	813	814	815	816	817

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Last AA of ORF	756	81	29	39	39	39	32	10	92	103
First AA of Secreted Portion	2	50		18	18	18	24		20	22
Last AA of Sig Pep	_	49		17	17	17	23		19	21
First Last AA AA of of Sig Sig Pep Pep	1	1	-	1	1	1	1	-	-	
AA SEQ ID NO: Y	3425	3426	3427	3428	3429	3430	3431	3432	3433	3434
5' NT of First AA of Signal Pep	2	20	141	307	307	307	230	169	21	138
5' NT of Start Codon		20	141	307	307	307	230	169	21	138
	3011	1445	1003	1838	1838	1838	1177	1568	1098	1122
5' NT of Clone Seq.	1		-	237	237	237	-	П	-	1
Total NT Seq.	3011	1445	1003	1901	1901	1901	1177	1731	1098	1122
NT SEQ ID NO: X	828	829	830	831	832	833	834	835	836	837
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203960 04/26/99	203960 04/26/99	203979 04/29/99	203979 04/29/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HFCA140	HFCAQ17	HFCBC16	HFCBL53	HFCBL53	HFCBL53	HFCBT29	HFCCZ31	HFCDN13	HFCDT67
Gene No.	818	819	820	821	822	823	824	825	826	827

r-		1		1	r			г	Υ	1
Last AA of ORF	30	93	46	161	11	57	89	ς,	27	34
First AA of Secreted Portion	16	16	19	22	:	30	17		18	20
Last AA of Sig Pep	15	15	18	21		29	16		17	19
First Last AA AA of of Of Sig Sig Pep Pep	1	-	1			1	1		-	1
AA SEQ ID NO: Y	3435	3436	3437	3438	3439	3440	3441	3442	3443	3444
5' NT of First AA of Signal Pep	628	164	238	43	6	385	228	542	136	93
5' NT of Start Codon	628	164	238	43	6	385			136	93
3' NT of Clone Seq.	829	1227	1513	650	3652	814	1059	934	066	896
5' NT of Clone Seq.	П	-			-	205		359	_	-
Total NT Seq.	829	1227	1513	650	3652	814	1059	896	066	896
NT SEQ ID NO:	838	839	840	841	842	843	844	845	846	847
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Lambda ZAP II
ATCC Deposit No.Z and Date	203960 04/26/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203957 04/26/99	PTA- 181 06/07/99	203960 04/26/99	PTA- 181 06/07/99	203960 04/26/99
cDNA Clone ID	HFCDY36	HFCEC45	HFCET43	HFEAG55	HFEAU63	HFEBA88	HFEBK75	HFEBO15	HFEBO17	HFFAE46
Gene No.	828	829	830	831	832	833	834	835	836	837

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Last AA of ORF	20	26	20	27	12	43	101	21	27	37
First AA of Secreted Portion	19	25	30	24		35	25	21	21	23
Last AA of Sig Pep	18	24	29	23		34	24	20	20	22
AA First SEQ AA ID of NO: Sig Y Pep	-		_	-	T		_	-	1	1
	3445	3446	3447	3448	3449	3450	3451	3452	3453	3454
5' NT of First AA of Signal	72	178	26	1833	347	507	259	183	30	302
5' NT of Start Codon	72	178	56	1833		507	259	183	30	302
3' NT of Clone Seq.	692	1134	1643	2298	1952	1076	561	1629	1018	892
5' NT 3' NT of of Clone Clone NT Seq. Seq.		-	-	1567	91	347	П	П	-	1
-	818	1134	1643	2298	1952	1076	561	1629	1018	892
NT SEQ ID NO:	848	846	850	851	852	853	854	855	856	857
Vector	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Other	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203918 04/08/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 1838 05/09/00	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HFFAH01	HFFAL70	HFFAV61	HFGAB50	HFGAE28	HFGAN63	HFHDN80	HFIAB78	HFIAD23	HFIAK06
Gene No.	838	839	840	841	842	843	844	845	846	847

Last AA of ORF	47	47	26	26	26	26	37	19	06	40
First AA of Secreted Portion	26	20	19	19	19	19	18	19	18	39
Last AA of Sig Pep	25	19	18	18	18	18	17	18	17	38
First Last AA AA of of of Sig Sig Pep Pep	-	-	-	-	-		-	-	-	-
AA SEQ ID NO:	3455	3456	3457	3458	3459	3460	3461	3462	3463	3464
5' NT of First AA of Signal	Pep 120	322	1611	1611 3458	1611	1611	280	268	2	1274 3464
5' NT of Start Codon	120	322	1611	1611	1611	1611	280	268		1274
3' NT of Clone Seq.	651	1270	3145	3145	3145	3145	1262	388	408	
S' NT 3' NT of of of Otatal Clone Clone NT Seq. Seq.	1	-	1366	1366 3145	1366	1366	-	-		1227 2967
_	651	1270	3145	3145	3195	3195	1262	388	408	867 3014
SEQ NO:	858	859	098	861	862	863	864	965	998	198
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203960	203960 04/26/99	203979 04/29/99	203979 04/29/99	203979 04/29/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99	203960 04/26/99
cDNA Clone ID	HFICH70	нғін057	HFIIK29	HFIIK29	HFIIK29	HFIIK29	нғпQ27	НЕПО64	HFIIZ61	HFIJD81
Gene No.	848	849	850	851	852	853	854	855	856	857

Last AA of ORF	35	32	34	74	24	38	59	9	31	35	36
First AA of Secreted Portion	22	28	28	24	12	28	33	29	19	19	23
AAA of Sig	21	27	27	23	11	27	32	28	18	18	22
AA First I SEQ AA DO DO Of NO: Sig Y	1	1	1	1	-	1	1	1	1	1	1
	3465	3466	3467	3468	3469	3470	3471	3472	3473	3474	3475
5' NT AA of SEQ First ID AA of NO: Signal Y Pep	82	306	87	51	404	227	774	86	69	89	243
5' NT of Start Codon	82	306	87	51		227	774	86	69	89	243
3' NT of Clone Seq.	1510	1207	839	1332	1978	979	1882	820	2461	1793	1005
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	6	22	-	-	171	Н	531	-	-	П	203
Total NT Seq.	1572	1207	839	1332	1978	626	1882	820	2485	1793	1005
NT SEQ ID NO:	898	698	028	871	872	873	874	875	928	877	878
Vector	pSport1										
ATCC Deposit No.Z and Date	203918 04/08/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	203918 04/08/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HFIJF44	HFITA02	HFITF80	HFTUK66	HFIUT21	HFIVB04	HFIXC39	HFIXC69	HFIXE39	HFIYP15	HFIZE10
Gene No.	828	829	098	861	862	863	864	865	998	867	898

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Last AA of ORF	99	24	37	35	30	111	32	113	172	172
First AA of Secreted Portion	56	24	25	25	17	11	20	29	10	10
	25	23	24	24	16	10	19	28	6	6
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	_	1	_	_	-	-	-		-
AA SEQ ID NO:	3476	3477	3478	3479	3480	3481	3482	3483	3484	3485
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	186	435	143	120	9	964	218	211	414	414
5' NT of Start Codon	186	435	143	120						
5' NT 3' NT of of Total Clone Clone NT Seq. Seq. Seq.	384	548	499	1289	1182	1648	1058	1332	2010	2023
5' NT of Clone Seq.	1	128		-	1		-	1		7
	384	548	499	1289	1182	1648	1058	1332	2010	2059
NT SEQ ID NO:	628	880	881	882	883	884	885	988	887	888
Vector	pSport1	pSport1	pSport1	Uni-ZAP XR						
ATCC Deposit No.Z and Date	203960 04/26/99	PTA- 181 06/07/99	203960 04/26/99	203979 04/29/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HFIZF51	HFIZK42	HFIZM89	HFKBA62	HFKBC47	HFKDX53	HFKEB14	HFKEG63	HFKES35	HFKES35
Gene No.	698	870	871	872	873	874	875	928	877	878

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Last AA of ORF	34	89	23	65	19	34	35	35	14	32
First AA of Secreted Portion	20	24	12	17	,	23	23	23		23
	61	23	11	16		22	22	22		22
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	-	1		_	-	_	1	-	1
AA SEQ ID NO: Y	3486	3487	3488	3489	3490	3491	3492	3493	3494	3495
5' NT of First AA of Signal Pep	229	382	1033	211	181	335	695	695	127	56
5' NT AA F 5' NT of SEQ of Start AA of NO: 3 Codon Signal Y I	229	382		211	181	335	695	695	127	56
S' NT 3' NT of of Clone Clone Seq.	1284	531	1980	2501	672	1942	2306	2306	779	715
S' NT 3' NT of Of Clone Seq. Seq.	1	69	289	1	1	_	360	360	1	
Total NT Seq.	1284	1288	1980	2501	672	1947	2311	2311	779	715
NT SEQ ID NO: X	688	068	891	892	893	894	\$68	968	268	868
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1
ATCC Deposit No.Z and Date	203960 04/26/99	PTA- 181 06/07/99	203960 04/26/99	203960 04/26/99	203979 04/29/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203960 04/26/99	203918 04/08/99
cDNA Clone ID	HFKEU17	HFKEV77	HFKFI15	HFKFI35	HFKFK49	HFKFV88	HFKFV88	HFKFV88	HFKFX64	HFOXD49
Gene No.	826	088	881	882	883	884	885	988	887	888

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Last AA of ORF	47	19	31	31	39	10	17	17	29	41
First AA of Secreted Portion	23		21	19	24		14	14		20
Last AA of Sig Pep	22		20	18	23		13	13		19
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep		1		1	-	1		1	1	1
AA SEQ ID NO: Y	3496	3497	3498	3499	3500	3501	3502	3503	3504	3505
5' NT of First AA of Signal		278	13	202	∞	199	347	347	205	230
5' NT of Start Codon		278	13	202	∞	199			205	230
3' NT of Clone Seq.	2053	396	916	1860	1490	783	1900	1900	732	802
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	-	2	1	1	1		1
	2053	396	916	1860	1490	783	1900	1900	732	802
NT SEQ ID NO:	668	006	901	902	903	904	905	906	907	806
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203979	203960	203960 04/26/99	PTA- 181 06/07/99	203960	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99	203960 04/26/99
cDNA Clone ID	HFOXE28	HFOYH74	HFOYP02	HFOYR24	HFOYR54	HFOZB26	HFPBF54	HFPBF54	HFPB193	HFPBJ64
Gene No.	688	068	891	892	893	894	895	968	897	868

Last AA of ORF	15	35	56	33	3	31	38	33	08	35	21
	 -	<i>c</i>	2	3		3	3	3	<u>~</u>	3	2
First AA of Secreted Portion		23	23	21		16	23	29	23	31	21
Last AA of Sig Pep		22	22	20		15	22	28	22	30	20
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	-	1			-	-		_	1		1
	3506	3507	3508	3509	3510	3511	3512	3513	3514	3515	3516
5' NT of of of Start AA of Codon Signal Pen	I.	56	194	57	101	12	292	277	88	238	401
	317	56	194	57		12	292	277	88	238	401
3' NT of Clone Seq.	846	1434	761	441	452	1699	1612	963	2210	1661	533
S' NT 3' NT of of Clone Clone Seq. Seq.		_	-	-		_	-	-	-	_	
Total NT Seq.	846	1434	761	441	452	1699	1612	963	2234	1661	533
SEQ NO:	606	910	911	912	913	914	915	916	917	918	616
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203979 04/29/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HFPBQ55	HFPCK22	HFPCM32	HFPCM36	HFPCS84	HFPCU47	HFPCY66	HFPDC65	HFPDE42	HFPDE88	HFPD025
Gene No.	668	006	901	905	903	904	905	906	206	806	606

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Last AA of	ORT	38	83	86	36	14	24	49	48	41	32	36
	Portion	25	30	35	20		16	35	44	36	27	20
Last AA of Sig		24	29	34	16		15	34	43	35	26	19
	rep	1	1	1	1	-	1-1		1	1	-	-
AA SEQ ID NO:	Ĭ	3517	3518	3519	3520	3521	3522	3523	3524	3525	3526	3527
5' NT AA of SEQ First ID AA of NO:	⊸ I	168	1455	64	150	218	12	132	100	130	212	284
5' NT of Start	Codon	168	1455	64	150		12	132	100	130	212	284
3' NT of Clone Seq.		2099	1837	922	1080	955	1164	1929	1444	878	793	1441
5' NT 3' NT of Olone Clone Seq. Seq.			1080	19	-	П	1	-		_	-	-
Total NT	Seq.	2099	1861	993	1080	955	1164	1929	1444	878	793	1441
		920	921	922	923	924	925	926	927	928	929	930
Vootor	Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR						
ATCC Deposit No.Z	and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203979 04/29/99	203960 04/26/99	203960	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA		HFPDP70	HFPDR39	HFPDX08	нгрер69	HFRAU40	HFRAY90	HFSAY91	HFSBC10	HFSBE94	HFTAN11	HFTAR27
Gene	NO.	910	911	912	913	914	915	916	617	918	616	920

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Last AA of ORF	56	23	136	33	33	70	∞	35	13	32	14
First AA of Secreted Portion	16	20	17	25	25	20		31		21	6
Last AA of Sig Pep	15	19	16	24	24	19		30		20	8
First Last AA AA of of Sig Sig Pep Pep	1	-		-	-	-	-		П		1
	3528	3529	3530	3531	3532	3533	3534	3535	3536	3537	3538
5' NT of First AA of Signal	242	249	94	49	49	363	250	206	1672	36	344
5' NT of Start Codon	242	249	94	49	49	363	250	206	1672	36	
	979	518	1830	1000	1077	1077	1309	910	2894	837	1377
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-	1	-	-	-	123	-	-	1572	_	1
Total NT Seq.	626	518	1830	1022	1077	1077	1309	910	2894	837	1377
SEQ NO:	931	932	933	934	935	936	937	938	939	940	941
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960	203979 04/29/99	203960 04/26/99	203979 04/29/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203959 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HFTAR30	HFTAS49	HFTBB50	HFTBL17	HFTBL17	HFTCF02	HFTC185	HFTCJ32	HFTC017	HFTCW07	HFTDF32
Gene No.	921	922	923	924	925	976	927	928	929	930	931

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Last AA of	ORF	31	187	25	21	27	9	93	12	13	39	62
First AA of Secreted	Portion	19	19	24		14		25		13	22	28
Last AA of Sig	Pep	18	18	23		13		24		12	21	27
	Pep	1	1	1	-			1	-	1	-	1
	Y	3539	3540	3541	3542	3543	3544	3545	3546	3547	3548	3549
5' NT of First AA of	Codon Signal Pep	617	139	216	375	2083	1198	462	46	962	146	203
5' NT of Start	Codon	617	139	216	375				46			203
		1319	1996	1200	1295	2163	1781	2151	1829	1581	1263	1347
5' NT 3' NT of of Clone Clone NT Seq. Seq.		242	321	-	84	249	-	62		731		-
Total NT	Seq.	1319	2014	1200	1295	2163	1781	2151	1829	1581	1263	1347
SEQ BD NO:	X	942	943	944	945	946	947	948	949	950	951	952
	Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript	pBluescript	pBluescript	pBluescript	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II
ATCC Deposit No.Z	and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203917 04/08/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
	Clone ID	HFTDF79	HFTDK11	HFTDU08	HFVGK67	HFVHD38	HFVHY57	HFVIC33	HFXAK32	HFXAK59	HFXBI64	HFXBL05
Gene	No.	932	933	934	935	936	937	938	939	940	941	942

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Last AA of	ORF	30	24	45	136	19	26	14	4	∞	36	21
First AA of Secreted	Portion ORF	28	18	29	25	18	10	14			16	19
Last AA of Sig		27	17	28	24	17	6	13			15	18
AA First Last SEQ AA AA ID of of NO: Sig Sig	Pep	1	-	-	-	1	-	-	1	1		1
	Y	3550	3551	3552	3553	3554	3555	3556	3557	3558	3559	3560
5' NT of First AA of	Signal Pep	232	263	52	68	179	275	278	49	48	237	159
5' NT of Start	Codon Signal Pep	232	263	52	68	179		278	49	48	237	159
3' NT of Clone Seq.		1277	1456	1728	498	502	1099	1757	1326	1237	909	1391
5' NT 3' NT of of Clone Clone Seq. Seq.		1	П	-	-	-	T	171	-			-
Total NT	Seq.	1277	1456	1728	498	502	1099	1757	1326	1237	1127	1391
SEQ BD NO:	X	953	954	955	926	957	958	959	096	961	962	963
	Vector	Lambda ZAP II										
ATCC Deposit No.Z	and Date	203960 04/26/99										
cDNA	Clone ID	HFXBM52	HFXBR58	HFXBV67	HFXBY20	HFXCB70	HFXCI42	HFXCL59	HFXCM22	HFXCN18	HFXCS53	HFXDB37
Gene	No.	943	944	945	946	947	948	646	026	951	952	953

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Last AA of	ORF	56	26	30	21	6	25	34	21	21	78	11
First AA of Secreted	Portion			61	20		24	19	11	11	34	
Last AA of Sig	Pep			18	19		23	18	10	10	33	
First Last AA AA of of Sig Sig	Pep	1		-	-	П	-		-	-	-	-
AA SEQ ID NO:	Y	3561	3562	3563	3564	3565	3566	3567	3568	3569	3570	3571
5' NT of First AA of	Signal Pep	204	186	11	71	308	258	79	338	1086	207	251
5' NT of Start	Codon Signal Pep	204	186	=	71		258	62			207	251
		1856	1558	1858	1760	588	1453	775	824	1298	1808	1349
5' NT 3' NT of of Clone Clone NT Seq. Seq.		Ţ	-	-		1	-	89	_	749	-	-
Total NT	Seq.	1856	1558	1858	1760	588	1453	775	824	1298	1808	1349
NT SEQ ID NO:	×	964	965	996	<i>L</i> 96	896	696	970	971	972	973	974
	Vector	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II							
ATCC Deposit No.Z	and Date	203960 04/26/99	203960	203959 04/26/99	203918 04/08/99							
cDNA	Clone ID	HFXDI32	HFXDJ43	HFXDL76	HFXDM75	HFXD018	HFXDP44	HFXDR08	HFXDR28	HFXDR28	HFXDR47	HFXDZ03
Gene	No.	954	955	926	156	856	959	096	961	362	963	964

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Last AA of ORF	34	28	16	73	91	32	39	29	62	92
First AA of Secreted Portion	20	22		25	18	24	17	25	23	40
Last AA of Sig Pep	61	21		24	17	23	16	24	22	39
First AA of Sig Pep	1			-	-		-	-	1	1
AA SEQ ID NO: Y	3572	3573	3574	3575	3576	3577	3578	3579	3580	3581
5' NT AA of SEQ First ID AA of NO: Signal Y Pep	112	59	267	42	125	158	240	70	217	302
5' NT of of of Start AA of Codon Signal Pep	112	59		42	125	158	240	70	217	302
3' NT of Clone Seq.	1953	1632	1363	1302	1230	361	1603	1647	1497	1566
5' NT 3' NT of Ol Clone Clone Seq.	1			-	-	-	-	-	-	1
Total NT Seq.	1953	1632	1363	1302	1230	361	1603	1647	1497	1566
NT SEQ ID NO: X	975	926	776	978	626	086	981	982	983	984
Vector	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 793 09/27/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HFXED33	HFXEE88	HFXGR32	HFXGT51	HFXGW16	HFXHC15	HFXHI33	HFXHL21	HFXHL83	HFXHM49
Gene No.	596	996	<i>L</i> 96	896	696	026	971	972	973	974

		ATCC		NT SEQ		S' NT 3' NT of		F	5° NT of	AA SEQ	First AA	Last	First	Last
Gene	cDNA	Deposit No.Z		A Ö	Total NT	Clone Clone Seq. Seq.	Clone Seq.	of Start	First AA of	ЭÖ	of Sig	of Sig	AA of Secreted	AA of
No.	Clone ID	and Date	Vector	×	Seq.		ı	Codon Signal Pep	Signal Pep	Y		Pep		ORF
576	нғхнм93	PTA- 181 06/07/99	Lambda ZAP II	586	1782	1	1782	307	307	3582		28	29	70
926	HFXHN89	203960 04/26/99	Lambda ZAP II	986	1406	1	1406	138	138	3583	-	23	24	34
277	HFXJB21	203960 04/26/99	Lambda ZAP II	282	1311	-	1311	232	232	3584	-	70	21	54
876	HFXJN93	203960 04/26/99	Lambda ZAP II	886	1742		1742	197	197	3585		23	24	4
626	HFXJS15	203959 04/26/99	Lambda ZAP II	686	1877	62	1877	356	356	3586	-			7
086	HFXJT53	203960 04/26/99	Lambda ZAP II	066	3013	105	3013	111	111	3587	-	39	40	361
981	HFXKG56	203960 04/26/99	Lambda ZAP II	991	992		992	273	273	3588		25	26	37
982	HFXKL60	PTA- 795 09/27/99	Lambda ZAP II	992	3138	200	3104	210	210	3589	-	30	31	33
983	HFXLG08	PTA- 793 09/27/99	Lambda ZAP II	993	1698		1698	306	306	3590	-	25	26	104
984	HFXLK91	203960 04/26/99	Lambda ZAP II	994	1848	1	1848	120	120	3591	-	20	21	399

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Last AA of ORF	23	39	61	48	71	7	09	71	30	32
First AA of Secreted Portion		33	17	32	26		25	24	21	24
Last AA of Sig Pep		32	16	31	25		24	23	20	23
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep	1	-	1		_	1	-	-	-	1
	3592	3593	3594	3595	3596	3597	3598	3599	3600	3601
5' NT of of Of Start AA of Codon Signal Pep	378	151	251	25	52	231	548	243	30	1005
5' NT of Start Codon	378	151	251	25	52		548	243	30	1005
5' NT 3' NT of of Clone Clone Seq.	740	1012	1693	1216	1191	1409	1838	695	989	2310
5' NT of Clone Seq.			1	-	-	1	454	27	1	816
Total NT Seq.	740	1015	1906	1216	1191	1418	1854	695	989	1004 2310
NT SEQ D NO:	995	966	766	866	666	1000	1001	1002	1003	1004
Vector	Lambda ZAP II	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203960 04/26/99	203959 04/26/99	PTA- 181 06/07/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203917 04/08/99	203960 04/26/99	203917 04/08/99
cDNA Clone ID	HFXLM32	HGBAX83	HGBBR29	HGBDL51	HGBDV35	HGBDX28	HGBGX31	НСВНЕ23	HGBHI15	HGCMW39
Gene No.	586	986	286	886	686	066	991	366	993	994

Last AA of	34 34	28	07	42	30	56	37	72	39	78	48
First AA of Secreted	rordon 19	28	20	16	10		19	37	18	21	42
Last AA of Sig	rep 18	77	/7	15	6		18	36	17	20	41
AA First Last SEQ AA AA ID of of of NO: Sig Sig		-		1	-	-	1	1	-	-	-
AA SEQ ID NO:	1 3602	3603	COOC	3604	3605	3606	3607	3608	3609	3610	3611
		691		161	91	191	546	39	310	254	268
5' NT of of of First Start AA of Signal	39	791	102	161	91		546	39			268
3' NT of Clone Seq.	774	614	110	849	762	778	1619	962	841	458	1537
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-		7	1	П	-	291			-	-
Total NT	жч. 774	614	017	849	762	778	1621	962	841	458	1537
NT SEQ ID NO:	1005	1006	7001	1007	1008	1009	1010 1621	1011	1012	1013	1014
Vector	vector Uni-ZAP XR	I Ini-ZAP XR		Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 1 1014
ATCC Deposit No.Z	allu Date 203960	203960	04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99	PTA- 181 06/07/99	203960 04/26/99	PTA- 181 06/07/99
cDNA	HGLAG32	HGLAH08	1	HGLAH86	HGLBC33	HGLBG15	HGLBM55	HGLDA95	HGLDB06	HGLDE15	HHBEI14
Gene	995	966	?	266	866	666	1000	1001	1002	1003	1004

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Last AA of ORF	21	36	54	116	32	46	42	36	51	106
First AA of Secreted Portion	19	10	20	35	15	20	26	21	.21	22
Last AA of Sig Pep	18	6	19	34	14	19	25	20	20	21
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	-		-		1	-	-		-	1
	3612	3613	3614	3615	3616	3617	3618	3619	3620	3621
		58	72	51	128	54	282	31	1334	70
5' NT of of of Start AA of Codon Signal	219	58	72	51	128	54	282	31	1334	70
3' NT of Clone Seq.	519	1734	1830	513	1030	1063	1749	138	1985	1576
5' NT 3' NT of of Clone Clone Seq. Seq.	1	-	∞	-			-	-	1320	1
Total NT Seq.	519	1734	1908	513	1030	1063	1749	138	1985	1576
SEQ NO NO	1015	1016	1017	1018		1020		1022	1023	1024
Vector	pCMVSport 1	pCMVSport 3.0 1016	pCMVSport 3.0 1017	pCMVSport 3.0 1018	pCMVSport 3.0 1019	pCMVSport 3.0 1020	pCMVSport 3.0 1021	pCMVSport 3.0 1022	pCMVSport 3.0 1023	pCMVSport 3.0 1024 1576
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203960 04/26/99		PTA- 795 09/27/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HHBGL33	HHEAW44	HHEBP28	HHECK41	HHECR10	HHEMC55	ннемм20	HHEMM80	HHEMP35	HHEMZ08
Gene No.	1005	1006	1007	1008	1009	1010	1011	1012	1013	1014

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Last AA of ORF	36	18	62	30	18	54	10	7	34	45	7
First AA of Secreted Portion	32		19	25	17	43			21	40	
Last AA of Sig Pep	31		18	24	16	42			20	39	
First Last AA AA of of Sig Sig Pep Pep	1	1	1	1	1	1	1	1	1	1	1
AA SEQ ID NO: Y	3622	3623	3624	3625	3626	3627	3628	3629	3630	3631	3632
5' NT of First AA of Signal Pep	187	206	5	143	87	790	344	1393	559	476	180
5' NT of Start Codon	187		5	143	87	790	344	1393	559	476	180
	1699	1126	1141	1580	2138	1154	1060	1973	2020	747	735
5' NT 3' NT of of Clone Clone NT Seq. Seq.	11	-	Т	Ţ	П	537	П	1017	520	_	1
Total NT Seq.	2238	1126	1141	1580	2138	2489	1060	3333	2020	747	735
NT SEQ ID NO:	1025	1026	1027		1029	1030	1031		1033	1034	1035
Vector	pCMVSport 3.0 1025	pCMVSport 3.0 1026	pCMVSport 3.0 1027	pCMVSport 3.0 1028	pCMVSport 3.0 1029	pCMVSport 3.0 1030	pCMVSport 3.0 1031	pCMVSport 3.0 1032	pCMVSport 3.0 1033	pCMVSport 3.0 1034	pCMVSport 3.0 1035
ATCC Deposit No.Z and Date	203960 04/26/99	203960	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960	203960 04/26/99	203960 04/26/99	203918 04/08/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HHENC17	HHENF95	HHENR74	HHENU33	HHENY07	HHEOK77	HHEPE72	HHEPE81	HHEPM64	HHEQI04	ннеоу60
Gene No.	1015	1016	1017	1018	1019	1020	1021	1022	1023	1024	1025

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Last AA of ORF	44	37	31	22	30	756	571		7	35
First Last AA of AA Secreted of Portion ORF	24	19	25	19	27	19	19			23
Last AA of Sig Pep	23	18	24	18	26	18	18	i		22
First Last AA AA of of Sig Sig Pep Pep	1	1	-	1			-	-		П
	3633	3634	3635	3636	3637	3638	5200	3639	3640	3641
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	131	89	161	24	183	120	715	1211	270	288
5' NT of Start Codon	131	89	161	24	183	120	715			
	1723	1054	1401	1447	1821	3168	2443	1302	1158	2046
5' NT of Clone Seq.	-	1		1	Н	П	-	_		232
Total NT Seq.	1723	1054	1401	1447	1821	3168	2443	1302	1158	2046
NT SEQ ID NO:	1036	1037	1038	1039	1040 1821	1041	2603	1042	1043	1044
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR 1044 2046
ATCC Deposit No.Z and Date	203960 04/26/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 181 06/07/99	PTA- 181 06/07/99	203960 04/26/99	203960 04/26/99	203979 04/29/99
cDNA Clone ID	HHFBA31	HHFCI81	HHFCN78	HHFCT95	HHFDN16	ннгев79	ннғев79	ннгес39	HHFEN34	HHFFZ01
Gene No.	1026	1027	1028	1029	1030	1031	1031	1032	1033	1034

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Last AA of	32	17	34	37	28	31	-	17	6	16	34
First AA of Secreted Portion				17	17	22					29
Last AA of Sig				16	16	21					28
First AA of Sig		1	1		-	-		-	П	П	-
AA SEQ ID NO:	3642	3643	3644	3645	3646	3647	3648	3649	3650	3651	3652
5' NT AA SEQ of First ID Start AA of NO:		89	19	213	17	200	231	223	363	314	121
5' NT of Start	62	89	19		17	200		223	363		121
5' NT 3' NT of Of Clone Clone Seq.	1590	1710	2764	1019	1097	724	859	1932	1302	545	1141
S' NT 3' NT of Of Clone Clone Seq. Seq.	1	-	1	-	-	_	-	-	201	_	1
Total NT Seg	ж. 1590	1711	2764	1019	1279	724	859	1932	1302	545	1141
NT SEQ ID NO:	1045	1046 1711	1047	1048	1049	1050	1051		1053	1054	1055
Vector	R	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Lambda ZAP II 1052	Lambda ZAP II 1053	Lambda ZAP II 1054	Lambda ZAP II 1055 1141
ATCC Deposit No.Z	203960	203960	203960	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HHFGI71	HHFGJ54	HHFGL38	HHFGR75	HHFGZ23	ннғнд26	HHFHM47	HHGAA76	HHGAD46	HHGAT09	HHGBC21
Gene	1035	1036	1037	1038	1039	1040	1041	1042	1043	1044	1045

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Last AA of ORF	46	126	99	49	69	38	19	84	36	20
First Last AA of AA Secreted of Portion ORF	13	27	19	22	23	16		27	20	16
First Last AA AA of of Sig Sig Pep Pep	12	26	18	21	22	15		26	19	15
First AA of Sig Pep	1	-	-		_	-	-		-	-
	3653	3654	3655	3656	3657	3658	3659	3660	3661	3662
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	36	25	220	122	234	137	9/9	21	73	113
5' NT of Start Codon	36	25	220		234	137	929	21	73	113
3' NT of Clone Seq.	655	798	1221	438	293	542	1026	1240	826	1174
S' NT 3' NT of Clone Clone Seq.		1		_	П	-	446	-	-	Н
Total NT Seq.	959	798	1221	438	442	542	1060	1240	826	1174
NT SEQ D NO:	1056	1057	1058	1059	1060	1061	1062		1064	1065
Vector	Lambda ZAP II 1056	Lambda ZAP II 1057	Lambda ZAP II	Lambda ZAP II 1059	Lambda ZAP II 1060	Lambda ZAP II 1061	Lambda ZAP II 1062 1060	Lambda ZAP II 1063	Lambda ZAP II 1064	Lambda ZAP II 1065
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99		PTA- 181 06/07/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HHGBF91	HHGBG63	HHGBV02	HHGBW55	HHGBX88	HHGCA26	HHGDA81	HHGDI12	HHGDR05	HHGDR92
Gene No.	1046	1047	1048	1049	1050	1051	1052	1053	1054	1055

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Last AA of ORF	27	32	39	36	11	26	10	31	39	42
First AA of Secreted Portion	18	30	28	35	47			24	15	30
Last AA of Sig Pep	17	29	27	34	46			23	14	29
First AA of Sig Pep	1	1	1	1	1	1	1		-	1
AA SEQ ID NO: Y	3663	3664	3665	3666	3667	3668	3669	3670	3671	3672
5' NT of First AA of Signal Pep	224	223	363	131	127	290	113	350	278	56
5' NT of Start Codon	224		363	131	127	290	113	350	278	56
5' NT 3' NT of Olone Clone Seq.	1502	814	1303	1522	1572	1625	1902	2054	1003	1832
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	1	1	_	1	1	1	1
	1502	814	1303	1522	1572	1631	1902	2054	1003	1832
NT SEQ ID NO: X	9901	1901	1068	1069	1070 1572	1071	1072 1902	1073 2054	1074 1003	1075 1832
Vector	Lambda ZAP II 1066 1502	Lambda ZAP II 1067	pBluescript SK- 1068	pBluescript SK- 1069 1522	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203959 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 791 09/27/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HHGDS56	HHGDW65	HHLBA86	HHNAC56	HHPBG90	ннРОЕ28	HHPDJ11	HHPDX86	HHPEA17	HHPEB61
Gene No.	1056	1057	1058	1059	1060	1061	1062	1063	1064	1065

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Last AA of ORF	80	37	30	38	5	11	28	34	99	33	34
First AA of Secreted Portion	28	22	27	29		26		17	38	29	15
Last AA of Sig Pep	<i>L</i> Z	21	26	28		25		16	37	28	14
First Last AA AA of of Sig Sig Pep Pep	1	1	1		-	1	1	П	1	1	-
AA SEQ ID NO: Y	3673	3674	3675	3676	3677	3678	3679	3680	3681	3682	3683
5' NT AA of SEQ First ID AA of NO: Signal Y	24	125	139	127	219	77	242	79	98	122	465
5' NT of of of Start AA of Codon Signal Pep		125	139	127		77	242	62	98	122	
5' NT 3' NT of of Clone Clone Seq. Seq.	2352	1050	746	2608	1067	2466	2549	1068	1546	1213	1250
5' NT 3' NT of of Clone Clone Seq. Seq.		П	-	П	П	-	-	П	1	18	330
Total NT Seq.	2352	1050	746	2608	1067	2466	2549	1068	1546	1392	1250
NT SEQ ID NO:	9201	1077	1078	1079	1080	1081	1082	1083	1084	1085	1086
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript	pBluescript	pBluescript	pBluescript
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	203960	203960	203960 04/26/99	203960	203960 04/26/99	203917 04/08/99
cDNA Clone ID	ННРFР26	HHPFS11	HHPFS15	HHPFS18	HHPGH34	HHPGU74	HHPGU87	HHPSD42	HHPSE03	HHPSE55	HHPSF70
Gene No.	1066	1067	1068	1069	1070	1071	1072	1073	1074	1075	1076

Last AA of ORF	29	39	30	38	9	22	28	16	74	27
First I AA of Secreted Portion C	27	39	56	19			17		14	18
Last AA of Sig Pep	76	38	25	18			16		13	17
First Last AA AA of of of Sig Sig Pep Pep	-	-	-	-	-	1	-	-		1
	3684	3685	3686	3687	3688	3689	3690	3691	3692	3693
5' NT of First AA of Signal Pep	167	65	23	248	333	151	78	161	310	432
5' NT of of of Start AA of Codon Signal Pep	167	99	23	248	333			161		432
3' NT of Clone Seq.	2107	1174	2029	1035	458	1610	1085	910	1654	1192
5' NT of Clone Seq.	-	-	П	162	-	_	Н	-	-	357
Total NT Seq.	2107	1174	1089 2029	1035	458	1610	1085	910	1654	1193
NT SEQ ID NO:	1087	1088	1089	1090 1035	1091	1092	1093	1094	1095	1096 1193
Vector	pBluescript	pBluescript	pBluescript	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203979 04/29/99	203960 04/26/99	PTA- 181 06/07/99	203960 04/26/99	203918 04/08/99	203979 04/29/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HHPSH74	HHPSL14	HHPSM40	HHPTF26	HHSAD31	HHSAE74	HHSAG62	HHSAK17	HHSBJ92	HHSBN84
Gene No.	1077	1078	1079	1080	1081	1082	1083	1084	1085	1086

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Last AA of ORF	25	40	36	433	38	39	28	31	17	20
First AA of Secreted Portion	∞	30	18	18	24	17	22	19	16	22
First Last AA AA of of Sig Sig Pep Pep	L	29	17	17	23	16	21	18	15	21
First AA of Sig Pep	1	-	-	-	-	1	1	1	-	1
AA SEQ ID NO: Y	3694	3695	3696	3697	3698	3699	3700	3701	3702	3703
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	275	112	175	46	22	40	591	145	847	55
	•	112	175	46	22	40	591	145	847	55
3' NT of Clone Seq.	983	847	282	1366	429	1721	1287	1288	1037	946
5' NT 3' NT of of Clone Clone Seq. Seq.	I			-	-	23	-	-	454	-
Total NT Seq.	983	847	282	2707	429	1721	1287	1290	1037	946
NT SEQ D NO:	1097	1098	1099	1100	1101	1102	1103	1104	1105	1106
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	ZAP Express
ATCC Deposit No.Z and Date	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	PTA- 1838 05/09/00	203979 04/29/99	203960 04/26/99	203960 04/26/99	203960 04/26/99	203960 04/26/99
cDNA Clone ID	HHSCL24	ННЅСО67	HHSCU12	HHSDB43	HHSDL07	HHSDX07	HHSFF54	HHSGB85	HHSGL84	ннтг.н79
Gene No.	1087	1088	1089	1090	1091	1092	1093	1094	1095	1096

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Last AA of ORF	65	34	43	55	24	53	35	11	28	38
First AA of Secreted Portion	32	31	24	28		24	23	,	20	23
Last AA of Sig Pep	31	30	23	27		23	22		19	22
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep		-	1	1		1	_		-	1
AA SEQ ID NO:	3704	3705	3706	3707	3708	3709	3710	3711	3712	3713
	Pep 130	140	124	104	47	779	331	73	944	597
5' NT of Start Codon	130	140	124	104	47	779	331	73	944	597
3' NT of Clone Seq.	1636	409	1652	1528	1790	2324	2913	424	1844	2124
5' NT of Clone Seq.		133	-	1	-	517		-	817	1
rotal NT Seq.	1636	409	1652	1528	1790	2324	2913	424	1844	2124
SEQ DD 7 NO:	1107	1108	1109	1110	1111	1112		1114	1115	1116 2124
Vector	Uni-ZAP XR	pBluescript	Other	Other	Other	Other	pBluescript SK- 1113	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203960	PTA- 181 06/07/99	PTA- 181 06/07/99	203960 04/26/99	203960 04/26/99	203960 04/26/99		203959 04/26/99	203918 04/08/99	203957 04/26/99
cDNA Clone ID	HIABC70	HIATG10	HIBCO70	HIBCR82	HIBDA41	HIBEC45	HILBW03	HISAE16	HISAG53	HISAN63
Gene No.	1097	1098	1099	1100	1101	1102	1103	1104	1105	1106

Last AA of ORF	9	36	25	38	33	32	406	[7	28	14
	99	3(2.	3	3;	33	40	317	5	1
First AA of Secreted Portion	30	33	18	25	29		2	2	27	
Last AA of Sig Pep	29	32	11	24	28		1	1	26	
First AA of Sig Pep	-	1	1	1	1	-	1	1	1	1
AA SEQ ID Y	3714	3715	3716	3717	3718	3719	3720	5201	3721	3722
5' NT of First AA of Signal Pen	<i>L</i> 96	169	1536 3716	257	82	315	480	698	193	311
5' NT of Start Codon	296	169	1536	257	82	315	480	698	193	
3' NT of Clone Seq.	2287	1058	2732	372	2043	1541	1699	1599	1796	1535
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	835		1309	-	П	116	1	1	1	
Total NT Seq.	2312	1058	2732	372	2043	1557	1699	1599	1796	1535
NT SEQ ID NO:	1117	1118	1119	1120	1121	1122	1123	2604	1124	
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pBluescript SK- 1124	pBluescript SK- 1125
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99							
cDNA Clone ID	HISAT78	HISBA38	HISBB66	HISCJ20	HISCK41	HISCO45	HISEJ52	HISEJ52	HJABC58	HJABG59
Gene No.	1107	1108	1109	1110	1111	1112	1113	1113	1114	1115

		ATCC		NT SEQ		S' NT 3' NT of		Ţ	5° NT of	• ,		Last AA	First	Last
Gene	cDNA	Deposit No.Z		ЭÃ	Total NT	Clone Clone Seq. Seq.		of Start	First ID AA of NO:	日党	of Sig	of Sig	AA of Secreted	AA of
No.	Clone ID	and Date	Vector	×	Seq.			Codon Signal Pep	Signal Pep	Y	Pep	Pep	Portion	ORF
1116	HJABR75	203957 04/26/99	pBluescript SK- 1126	1126	1328	П	1328	324	324	3723	-	18	19	37
1117	HJABS31	203957 04/26/99	pBluescript SK- 1127		1232	42	1232		-	3724	-	-	2	410
1118	HJABT12	203957 04/26/99	pBluescript SK- 1128	1128	557	-	557	30	30	3725	_	16	17	37
1119	HJACE25	203957 04/26/99	pBluescript SK- 1129	1129	1320	_	1320	322	322	3726	-			26
1120	HJACK21	203957 04/26/99	pBluescript SK- 1130	1130	1271	-	1271	139	139	3727	-	21	22	35
1121	HJBCG74	203957 04/26/99	pBluescript SK- 1131		2455	12	2428	33	33	3728	-	30	31	224
1122	HJBC021	203957 04/26/99	pBluescript SK- 1132	1132	587		553	289	289	3729	-			11
1123	НЈВСО40	203957 04/26/99	pBluescript SK- 1133	1133	1069	1-	1069	161	161	3730	-	20	21	51
1124	HJBDM36	203957 04/26/99	pBluescript SK- 1134 2777	1134	2777	1716 2760		1954	1954 3731	3731	-	18	19	40
1125	HJMAF30	203957 04/26/99	pCMVSport 3.0 1135	1135	603	-	603	193	193	3732		24	25	34
1126	HJMAM72	203957 04/26/99	pCMVSport 3.0 1136	1136	403	-	403	300	300	3733	-	17	18	25

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Last AA of ORF	45	36	36	36	70	45	29	58	33	33	290
First AA of Secreted Portion	56	26	26	26	25	43	16	19	18	17	51
Last AA of Sig Pep	25	25	25	25	24	42	15	18	17	16	50
First AA of Sig Pep	1	1	1	1		1	1	1	-		1
	3734	3735	3736	3737	3738	3739	3740	3741	3742	3743	3744
5' NT AA of SEQ of First ID Start AA of NO: Codon Signal Y	1441	1376	2308	2308	42	58	145	105	81	25	99
5' NT of Start Codon	1441	1376	2308	2308	42	58	145	105	81	25	99
3' NT of Clone Seq.	2915	3021	3953	3953	658	633	275	1439	1020	1076	1109
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1393	1305	2237	2237	-	1	-	-	Н	П	1
_	2968	3021	3953	3953	859	633	275	1439	1020	1076	1109
NT SEQ B NO:	1137	1138	1139	1140	1141	1142	1143	1144	1145	1146	1147 1109
Vector	pCMVSport 3.0 1137	pCMVSport 3.0 1138	pCMVSport 3.0 1139	pCMVSport 3.0 1140	pCMVSport 3.0 1141	pCMVSport 3.0 1142	pCMVSport 3.0 1143	pCMVSport 3.0 1144	pCMVSport 3.0 1145 1020	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203979 04/29/99	203957 04/26/99	203957 04/26/99	203957 04/26/99							
cDNA Clone ID	HJMAZ60	HJMBB20	HJMBB20	HJMBB20	HJMBK59	HJMBP01	HJMBQ17	НЈМВW62	HJMBX54	HJPAF69	HJPAQ19
Gene No.	1127	1128	1129	1130	1131	1132	1133	1134	1135	1136	1137

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Last AA of ORF	24	25	73	55	27	33	<i>L</i> 9	7	144	18
First AA of Secreted Portion		24	28	46		20	12		23	13
Last AA of Sig Pep		23	27	45		19	11		22	12
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1		1		-	1	-	П	П
	3745	3746	3747	3748	3749	3750	3751	3752	3753	3754
5' NT of of Strat AA of Codon Signal Pep	832	89	70	370	188	134	320	642	269	213
	832	89	. 70	370	188	134	320		269	
3' NT of Clone Seq.	1940	538	1036	936	905	1044	1417	1377	905	1813
5' NT 3' NT of of Clone Clone NT Seq. Seq.	599	1	1	234		-	78	324	-	145
Total NT Seq.	1963	808	1036	938	905	1044	1417	1377	905	1888
NT SEQ D NO:	1148	1149	1150	1151	1152	1153	1154 1417		1156	1157
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR 1150	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 2.0 1155	pCMVSport 2.0 1156	pCMVSport 2.0 1157
ATCC Deposit No.Z and Date	203979 04/29/99	203957 04/26/99	203957 04/26/99	203959 04/26/99	203957 04/26/99	203957 04/26/99	203918 04/08/99		203957 04/26/99	203917 04/08/99
cDNA Clone ID	HJPAZ35	HJPBI77	HJPBN96	HJPBU47	НЈРСQ19	HJPDJ08	HJPDK61	HKABI53	HKABN63	HKACA25
Gene No.	1138	1139	1140	1141	1142	1143	1144	1145	1146	1147

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Last AA of ORF	96	15	24	148	18	14	36	25	377	175
First AA of Secreted Portion	25			26			20	21	2	20
Last AA of Sig Pep	24			25			61	20	1	19
AA First SEQ AA ID of NO: Sig Y Pep	1	1	1	1	1		1	-	-	1
AA SEQ ID NO: Y	3755	3756	3757	3758	5202	3759	3760	3761	3762	3763
5' NT of First AA of Signal Pep	290	530	146	125	243	335	462	62	154	78
5' NT of Start Codon	290	530		125	243	335	462	62	154	78
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1898	1941	906	4597	2175	558	1442	1228	2241	1577
5' NT 3' NT of Of Clone Seq. Seq.	1	466	1	802	1	206	82	1	1	1
	1899	1987	906	4597	2175	558	1442	1228	2241	1577
NT SEQ ID NO:	1158	1159	1160	1161	2605	1162	1163	1164	1165	1166
Vector	pCMVSport 2.0	pCMVSport 2.0 1159 1987	pCMVSport 2.0 1160	pCMVSport 2.0 1161	pCMVSport 2.0 2605 2175	pCMVSport 2.0 1162	pCMVSport 2.0 1163 1442	pCMVSport 2.0 1164 1228	pCMVSport 2.0 1165	pCMVSport 2.0 1166 1577
ATCC Deposit No.Z and Date	PTA- 795 09/27/99	203957 04/26/99	203957 04/26/99	203918 04/08/99	203918 04/08/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HKACO64	HKACP50	HKACX90	HKADI27	HKADI27	HKADN26	HKADP79	HKADT55	HKAEK58	HKAEK72
Gene No.	1148	1149	1150	1151	1151	1152	1153	1154	1155	1156

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Last AA of ORF	31	59	48	11	417	38	19	30	31	37	44
First AA of Secreted Portion	25	26	22		2	27	16	30	21	17	23
Last AA of Sig Pep	24	25	21		-	26	15	29	70	16	22
First AA of Sig Pep	1	1		-	-	-		-			1
	3764	3765	3766	3767	3768	3769	3770	3771	3772	3773	3774
5' NT AA of SEQ First ID AA of NO: Signal Y Pep	145	487	462	73	1	33	114	142	290	151	762
5' NT of Start Codon	145	487	462	73		33	114	142	290	151	762
3' NT of Clone Seq.	2110	1791	1349	1273	1468	1176	1779	1473	622	1332	2129
5' NT 3' NT of Ol Clone Seq. Seq.	1	315	-	29			-	П	1		1
Total NT Seq.	2110	1825	1349	1273	1468	1176	1779	1174 1473	779	1332	1177 2129
NT SEQ ID NO:	1167	1168	1169	1170	1171	1172	1173 1779	1174	1175	1176	1177
Vector	pCMVSport 2.0 1167 2110	pCMVSport 2.0 1168	pCMVSport 2.0 1169	pCMVSport 2.0 1170	pCMVSport 2.0 1171 1468	pBluescript	ZAP Express	ZAP Express	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203918 04/08/99	203979 04/29/99	203979 04/29/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HKAFJ47	НКАГО41	HKAHH71	HKAJA95	HKAKU90	HKCSZ54	HKFAA15	HKFBB08	HKGAG59	HKGAJ81	HKGAK45
Gene No.	1157	1158	1159	1160	1161	1162	1163	1164	1165	1166	1167

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Last AA of ORF	33	4	25	38	39	32	25	45	42	29
First AA of Secreted Portion	33	31	23	27	22	17	19	42	26	22
First Last AA AA of of Sig Sig Pep Pep	32	30	22	26	21	16	18	41	25	21
	-	-	-	П	-	-	-	1	-	-
AA SEQ ID NO: Y	3775	3776	3777	3778	3779	3780	3781	3782	3783	3784
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	i	1155	379	109	43	301	225	351	36	136
	213	1155		109	43	301	225	351	36	136
3' NT of Clone Seq.	2332	1907	1639	1858	1036	849	1759	2220	2702	1410
5' NT of Clone Seq.	1	73	102	1	-	-	-	-	П	1
Total NT Seq.	2332	1907	1639	1858	1036	849	1759	2220	2702	1785
SEQ NO:	1178	1179	1180	1181	1182	1183	1184	1185	1186 2702	1187
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203979 04/29/99	203918 04/08/99	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 795 09/27/99	203957 04/26/99
cDNA Clone ID	HKGAP04	HKGAP57	HKGAW41	HKGBA21	HKGBC33	HKGBC73	HKGBF61	HKGBH54	HKGBP52	HKGCE23
Gene No.	1168	1169	1170	1171	1172	1173	1174	1175	1176	1177

Last AA of ORF	40	102	102	41	57	122	9	40	55	30
First I AA of Secreted Portion C	33	51	51		23	50	61	21	23	19
Last AA of Sig Pep	32	20	20		22	49	18	20	22	18
First AA of Sig	-	-	-	-	-		-	1	-	1
AA SEQ ID NO: Y	3785	3786	3787	3788	3789	3790	3791	3792	3793	3794
	44	308	316	303	329	302	145	26	751	235
5' NT of Start Codon	44	308	316	303	329		145	26	751	235
S' NT 3' NT of of Clone Clone Seq.	1162	1024	1040	1103	1658	1167	1671	909	1696	1994
5' NT of Clone Seq.	1	1	1		-		-	-	732	1
Total NT Seq.	1162	1024	2191	1103	1658	1167	1671	506	1721	1994
NT SEQ ID NO:	1188	1189	1190	1191	1192	1193	1194	1195		1197
Vector	pSport1	Lambda ZAP II	Lambda ZAP II 1196	pBluescript						
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HKGCE62	HKGCK41	HKGCK41	HKGCN96	HKGCX05	HKGDA95	HKGD012	HKIME53	HKIMG23	HKIXB73
Gene No.	1178	1179	1180	1181	1182	1183	1184	1185	1186	1187

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Last AA of ORF	38	46	31	32	17	33	30	15	294	40	32
First AA of Secreted Portion	27	24	31	23	15	25	18	14	30	31	19
	97	23	30	22	14	24	17	13	29	30	18
First Last AA AA of of Sig Sig Pep Pep	1	-	-	-	1	-		-	1	1	-
AA SEQ ID NO: Y	3795	3796	3797	3798	3799	3800	3801	3802	3803	3804	3805
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	133	10	183	196	396	59	203	287	329	87	494
5' NT of Start Codon	133		183	196		59	203	287	329	87	494
3' NT of Clone Seq.	443	1560	463	477	289	1877	782	1003	1692	1274	1601
5' NT of Clone Seq.		-	-	П	-		-	-	1		223
Total NT Seq.	443	1560	463	477	289	1877	782	1003	1692	1274	1601
NT SEQ ID NO:	1198	1199	1200	1201	1202	1203	1204	1205	1206	1207	1208
Vector	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript
ATCC Deposit No.Z and Date	2039 <i>57</i> 04/26/99	203957 04/26/99	203979 04/29/99	203957 04/26/99	203957 04/26/99						
cDNA Clone ID	HKIXD68	HKIXR91	HKIXS19	HKIXW45	HKIYU90	HKMLB81	HKMLF77	HKMLM32	HKMLR17	HKMLT89	HKMLV05
Gene No.	1188	1189	1190	1191	1192	1193	1194	1195	1196	1197	1198

Last AA of ORF	31	37	33	26	21	55	12	25	37	23
	(4.)	(T)	G.)	C4	(4	4.)			4.1	.,
First AA of Secreted Portion	27	17	21	6		30		12	32	18
Last AA of Sig Pep	26	16	20	8		29		11	31	17
First Last AA AA of of Sig Sig Pep	1	1	1	_	1	1	1	-	1	-
	3806	3807	3808	3809	3810	3811	3812	3813	3814	3815
	251	1109 3807	150	312	361	228	126	334	181	2010 3815
5' NT of of of Start AA of Codon Signal Pep	251	1109	150		361	228			181	2010
	992	3230	2070	1259	1905	1147	866	810	436	3714
5' NT of Clone Seq.	1	006	-	-	-	101	П	31	T	1908
Total NT Seq.	992	3237	2070	1259	1905	1147	866	810	436	1218 3714
NT SEQ ID NO:	1209	1210	1211	1212	1213	1214	1215	1216	1217	1218
Vector	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	Uni-ZAP XR	Uni-ZAP XR	pBluescript	Uni-ZAP XR
ATCC Deposit No.Z and Date	203957 04/26/99	203917 04/08/99	PTA- 181 06/07/99	203979 04/29/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HKMLV25	HKMMB79	HKMMC69	HKMMD91	HKMMP90	HKMMU76	HKPAC10	HKPAC50	HKPMA08	HKTAC18
Gene No.	1199	1200	1201	1202	1203	1204	1205	1206	1207	1208

Last AA of ORF	18	383	44	37	29	123	40	74	43	<i>L</i> 9
First AA of Secreted Portion		28	17	20		21	23	51	22	20
Last AA of Sig Pep		27	16	19		20	22	50	21	19
First Last AA AA of of Sig Sig Pep Pep	1	1		-		-	1	1		1
AA SEQ ID NO: Y	3816	3817	3818	3819	3820	3821	3822	3823	3824	3825
5' NT of First AA of Signal Pep	1030	145	41	281	38	372	1176	147	759	169
5' NT of Start Codon	1030				38		1176	147	759	169
	1219	1476	475	2708	1314	1022	2820	787	2480	779
S' NT 3' NT of of Clone Clone Seq. Seq.	1017	1	1	77	1		888	1	496	94
Total NT Seq.	1263	1220 1476	475	1222 2708	1314	1022	2820	787	2638	787
NT SEQ ID NO:	1219	1220	1221	1222	1223	1224	1225	1226	1227	1228
Vector	pBluescript	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 3.0 1223	pCMVSport 3.0 1224 1022	pCMVSport 3.0 1225	pCMVSport 3.0 1226	pCMVSport 3.0 1227	pCMVSport 3.0 1228
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 1838 05/09/00	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99	PTA- 181 06/07/99
cDNA Clone ID	HL1SA89	HL2AB60	HL3AE69	HL3AF32	HLDAV70	HLDBL62	HLDBV18	HLDBV54	HLDCR26	HLDDM27
Gene No.	1209	1210	1211	1212	1213	1214	1215	1216	1217	1218

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Last AA of ORF	<i>L</i> 9	129	36	31	46	55	51	51	4	45
First AA of Secreted Portion	20	24	30	26	27	26	21	37	23	21
Last AA of Sig Pep	19	23	29	25	26	25	20	36	22	20
First Last AA AA of of Sig Sig Pep Pep	1	-		1	-	-	-	-	-	
AA SEQ ID NO: Y	3826	3827	3828	3829	3830	3831	3832	3833	3834	3835
5' NT AA of SEQ of First ID Start AA of NO: Codon Signal Y Pep	177	197	12	106	193	238	161	28	34	46
5' NT of Start Codon	177	197	12	106	193	238	161	28	34	46
5' NT 3' NT of Olone Clone Seq.	788	1686	936	869	903	1971	1086	559	1425	2324
5' NT of Clone Seq.	102	175	-	1		85		-		1
Total NT Seq.	662	1726	936	869	903	1971	1086	559	1425	2324
NT SEQ D NO:	1229	1230	1231	1232	1233			1236	1237	1238 2324
Vector	pCMVSport 3.0 1229	pCMVSport 3.0 1230	pCMVSport 3.0 1231	pCMVSport 3.0 1232	pCMVSport 3.0 1233	pCMVSport 3.0 1234	pCMVSport 3.0 1235	pSport1	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203917 04/08/99
cDNA Clone ID	HLDDM27	HLDNF18		HLDOD77	HLDOL74	HLDPB24	HLDRU08	HLDXF43	HLEAA10	HLEAA24
Gene No.	1219	1220	1221	1222	1223	1224	1225	1226	1227	1228

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Last AA of ORF	40	40	343	55	52	51	16	48	10	-
First AA of Secreted Portion	23	23	2	22	12	15		16		
	22	22	-	21	11	14		15		
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1		-	-	-		-	1
• • • • • • • • • • • • • • • • • • • •	3836	3837	3838	3839	3840	3841	3842	3843	3844	3845
5' NT of of Of Start AA of Codon Signal Pep		17	73	6	423	140	157	237	182	63
5' NT of Start Codon	17	17	73	6	423	140	157	237	182	
3' NT of Clone Seq.	2032	2033	4023	1674	878	1134	1260	1818	2154	947
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.			2309	-		-	-	_	-	1
Total NT Seq.	2041	1240 2054	4038	1674	878	1134	1260	1818	2154	947
NT SEQ D NO:	1239	1240	1241	1242	1243	1244 1134	1245	1246 1818	1247	1248
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203957 04/26/99	203957 04/26/99	PTA- 793 09/27/99	203918 04/08/99
cDNA Clone ID	HLHAE14	HLHAE14	HLHBS54	HLHCB33	HLHCF14	HLHCG24	нгнсн20	HLHCN51	нгнст96	нгнрс33
Gene No.	1229	1230	1231	1232	1233	1234	1235	1236	1237	1238

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Last AA of ORF	32	21	7	38	38	38	38	41	15	09	61
First AA of Secreted Portion	15	18		21	21	21	21	12		19	24
Last AA of Sig Pep	14	17		20	20	20	20	11		18	23
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	1	·	-
AA SEQ ID NO: Y	3846	3847	3848	3849	3850	3851	3852	3853	3854	3855	3856
	515	73	94	432	433	433	433	282	157	153	1260
5' NT of of of Start AA of Codon Signal Pep	515	73		432	433	433	433		157	153	1260
	808	839	971	2332	2502	2556	2127	1105	1274	1491	2270
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	363	364	364	364	-	1		1107
Total NT Seq.	808	839	971	2351	2516	2556	2127	1256 1105	1274	1491	1259 3045
NT SEQ D NO:	1249	1250	1251	1252	1253 2516	1254	1255 2127	1256	1257 1274	1258	1259
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918	203918 04/08/99	203979	203979 04/29/99	203979 04/29/99	203979	203979 04/29/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203979 04/29/99
cDNA Clone ID	HLHDF92	HLHDJ05	HLHDL37	697ДНТН	69ТДНТН	HLHDL69	69ТОНТН	HLHDM38	HLHDR92	HLHDY94	HLHEE27
Gene No.	1239	1240	1241	1242	1243	1244	1245	1246	1247	1248	1249

Last AA of ORF	17	146	54	26	31	23	62	26	37	43
First AA of Secreted Portion (16	21	23	27	50	6 .	22	33	20
Last AA of Sig Pep		15	70	22	26	19	∞	21	32	19
First AA of Sig	-		-	1	-	—	1	-	1	1
AA SEQ ID NO:	3857	3858	3859	3860	3861	3862	3863	3864	3865	3866
	270	88	23	220	133	461	1368 3863	39	162	161
5' NT of of of Start AA of Codon Signal Pep	270	68	23	220	133	461		39	162	161
	880	1154	1124	816	1232	854	2092	1352	1658	774
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	-	-		-	1	1	1	П	_
Total NT Seq.	880	1154	1124	816	1232	854	2092	1352	1268 1658	774
SEQ BD NO:	1260	1261	1262	1263	1264 1232	1265	1266 2092	1267	1268	1269
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript	pBluescript	pBluescript	pBluescript
ATCC Deposit No.Z and Date	203957 04/26/99	PTA- 791 09/27/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203979 04/29/99
cDNA Clone ID	HLHEE38	HLHEI72	HLHEX62	HLHFK59	HLHFP09	HLHGG78	HLHSG15	HLHSQ35	HLHTB92	HLHTP55
Gene No.	1250	1251	1252	1253	1254	1255	1256	1257	1258	1259

Last AA of ORF	07	33	23	28	28	32	46	24	32	55	6
First Last AA of AA Secreted of Portion ORF			22	17	17	25	44	6	29	28	
Last AA of Sig S Pep	19	22	21	16	16	24	43	∞	28	27	
First AA of Sig Pep	1	_	1	1	1	1	1	_	1	1	-
AA SEQ ID NO: Y	3867	3868	3869	3870	3871	3872	3873	3874	3875	3876	3877
	127	198	467	416	38	282	66	237	249	160	63
5' NT of of of Start AA of Codon Signal Pep	127	198	467	416	38	282	66	237	249		63
	411	779	638	1055	1161	1681	8/9	610	1264	942	1522
5' NT 3' NT of of Clone Seq.	1	26	202	_	-		1	1	1	1	П
Total NT Seq.	411	622	638	1055	1161	1681	829	610	1264	942	1522
SEQ NO:	1270	1271	1272	1273	1274 1161	1275	1276	1277	1278	1279	1280 1522
Vector	pCMVSport 1	pCMVSport 1	pCMVSport 1 1272	pCMVSport 1	pCMVSport 1	pCMVSport 1	pCMVSport 1 1276	pCMVSport 1	Lambda ZAP II 1278	Lambda ZAP II 1279	Uni-ZAP XR
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HL.IBD74	HLIBE41	HLIBO16	HLJB122	HLJEE16	HLLAX64	HLLAX95	HLLCD67	HLMBX89	HLMBZ14	HLMCT51
Gene No.	1260	1261	1262	1263	1264	1265	1266	1267	1268	1269	1270

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Last AA of ORF	50	34	57	72	15	23	59	98	19	33
Last AA First of AA of Sig Secreted Pep Portion	44	22	17	22		17	10	45		18
Last AA of Sig Pep	43	21	16	21		16	6	44		17
First AA of Sig Pep	-	1	1	1	1	_1	1		1	
	3878	3879	3880	3881	3882	3883	3884	3885	3886	3887
f. I	Pep 195	183	7.1	21	268	185	520	141	21	271
5' NT of Start Codon	195		71	21	268	185		141	21	271
3' NT of Clone Seq.	1446	1193	621	1059	290	965	1175	1340	929	927
5' NT of Clone Seq.	-	-	-	П	П	-	236	1	1	
Total NT Seq.	1446	1193	921	1059	590	965	1175	1340	656	927
SEQ SEQ NO:	1281	1282	1283	1284 1059	1285	1286	1287	1288	1289	1290
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Lambda ZAP II	Lambda ZAP II 1286	Lambda ZAP II 1287	Lambda ZAP II 1288	Lambda ZAP II 1289	Lambda ZAP II 1290
ATCC Deposit No.Z and Date	203957	203957 04/26/99	PTA- 181 06/07/99	203957 04/26/99						
cDNA Clone ID	HLMCT95	HLMDD65	HLMDH01	HLMDU23	HLMFB62	HLMFG52	HLMFU53	HLMHG68	HLMHN06	HLMHS15
Gene No.	1271	1272	1273	1274	1275	1276	1277	1278	1279	1280

Last AA 1 of ORF	48	70	86	21	10	38	59	141	89	16
First AA of Secreted Portion	21		26	70		15	70	16	45	
Last AA of Sig Pep	20		25	19		14	19	15	44	
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	-	1	-	1	1	1	1	1		
AA SEQ ID NO: Y	3888	3889	3890	3891	3892	3893	3894	3895	3896	3897
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y		100	38	137	131	295	445	142	28	268
5' NT of Start Codon	l	100	38	137	131	295	445		28	268
3' NT of Clone Seq.	1635	1246	358	622	446	412	1006	6981	9/9	1001
for NT 3' NT of of Of Clone Clone NT Seq. Seq.	-	1	1	1	1	38	127	1		-
Total NT Seq.	1635	1246	358	622	446	445	1006	1369	9/9	1061
SEQ BD NO:		1292	1293	1294	1295	1296	1297	1298	1299	1300
Vector	Lambda ZAP II	Lambda ZAP II 1292	Lambda ZAP II	Lambda ZAP II 1294	Lambda ZAP II 1295	Lambda ZAP II 1296	Lambda ZAP II 1297 1006	Lambda ZAP II	Lambda ZAP II 1299	Lambda ZAP II 1300 1061
ATCC Deposit No.Z and Date	203957 04/26/99		203957 04/26/99			203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99	203957 04/26/99	203957
cDNA Clone ID	HLMIM84	HLMIN52	HLMIQ83	HLMIW76	HLMMA65	HLMMT12	HLMNA19	HLQAD72	HLQAM30	HLQAM59
Gene No.	1281	1282	1283	1284	1285	1286	1287	1288	1289	1290

Last AA of ORF	59	32	15	72	37	25	8	99	70	30	20
First L AA of A Secreted O Portion O		23		31	16	25	19	23	19	56	
Last AA of A AS Sig Se Pep Pep	56	22		30	15	24	18	22	18	25	
	_	_		-	-	1	1		1		-
	3898	3899	3900	3901	3902	3903	3904	3905	3906	3907	3908
	268	95	810	120	377	327	95	115	181	113	246
5' NT of of of Start AA of Codon Signal Pep	995	95	810	120		327	95	115	181	113	246
	2046	577	1349	1026	1103	1173	845	1781	919	495	1483
5' NT 3' NT of Ol Clone Seq. Seq.	442	1	762	-	-	-	1	1	144	-	-
Total NT Seq.	2046	577	2108	1026	1103	1421	845	1781	919	495	1483
SEQ NO:	1301	1302				1306	1307	1308	1309	1310	1311
Vector	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II 1303	Lambda ZAP II 1304	Lambda ZAP II 1305	Lambda ZAP II	Lambda ZAP II	Lambda ZAP II 1308	Lambda ZAP II 1309	pSport1	Uni-ZAP XR
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203959 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	нцовв23	нговгоз	HLQBX64	HLQCY09	HLQCZ43	HLQCZ80	HLQDK45	HLQDM47	нгори77	HLSAD72	HLTCJ67
Gene No.	1291	1292	1293	1294	1295	1296	1297	1298	1299	1300	1301

Last AA of ORF	18	42	50	37	37	38	34	140	11	33	79
First AA of Secreted Portion		30	24	26		29	14	24			44
Last AA of Sig Pep		29	23	25	25	28	13	23			43
First Last AA AA of of Sig Sig Pep Pep	1	1	1		1	-	1	1	1	1	1
	3909	3910	3911	3912	3913	3914	3915	3916	3917	3918	3919
5' NT of First AA of Signal Pep	66	758	66	120	120	520	127	761	22	53	71
5' NT of Start Codon	66	758	66			520			25	53	71
3' NT of Clone Seq.	1332	1672	947	1474	1474	1982	2689	1573	1986	1993	1469
5' NT of Clone Seq.	1	265	1		-	1	-	648	1		1
Total NT Seq.	1332	1676	947	1744	1744	1982	2689	1573	1986	1993	1469
NT SEQ D NO:	1312	1313	1314	1315	1316 1744	1317	1318 2689	1319	1320	1321	1322 1469
Vector	Uni-ZAP XR										
ATCC Deposit No.Z and Date	203957 04/26/99	203959 04/26/99	203957 04/26/99								
cDNA Clone ID	HLTCM28	HLTC022	HLTDA14	HLTDC26	HLTDC26	HLTDI20	HLTDI65	HLTDK30	HLTDL37	HLTDU35	HLTDX04
Gene No.	1302	1303	1304	1305	1306	1307	1308	1309	1310	1311	1312

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Last AA of	ORF	31	1		23	35		6		22		3	37	5	43		52
First AA of Secreted	Portion					17				19			18	2	26		10
	Pep					16	l			18			17	•	25		6
	Pep	1	1		1	-		1		1		1	1		-		1
	Y	3920	3921		3922	3923		3924		3925		3926	3927	1	3928		3929
5' NT of First AA of	Codon Signal Pep	88	210		122	251		218		229		257	1117		63		9/9
5' NT of Start	Codon	88			122	251		218		229			1117	•	63		
3' NT of Clone Seq.		1254	1827		1514	1535	 	2583		1290		1503	7287) 	2929		1203
5' NT 3' NT of of Clone Clone NT Seq. Seq.		1	_		-	-	ı	-		1		1	-	1	-		
Total NT	Seq.	1254	1827		1514	1535		3051		1290		1503	2289) 	2929		1203
. ~ ::	×	1323	1324		1325	1326 1535		1327		1328		1329	1330))			1332
	Vector	Uni-ZAP XR	Uni-ZAP XR		Uni-ZAP XR	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-Zap XR	nCMVSnort 3 0 1330 2289		pCMVSport 3.0 1331		pCMVSport 3.0 1332
ATCC Deposit No.Z	and Date	203957 04/26/99	PTA-	181 06/07/99	203957	203957	04/26/99	203959	04/26/99	203957	04/26/99	203957	04/20/22 PTA-	181	PTA-	09/27/99	203957 04/26/99
	Clone ID	нстен84	HLTEL39		HLTEN11	HLTEW52		HLTEZ36		HLTGG14		HLUAF94	HI WAH33		HLWA011		HLWAW73
Gene	No.	1313	1314		1315	1316		1317		1318		1319	1320) 	1321		1322

Last AA of ORF	39	72	11	31	9	32	33	53	26	22	7
First AA of Secreted Portion	29	48		31		18	<i>L</i> 1	44	2		
Last AA of Sig Pep	28	47		30		17	91	43	1		
First AA of Sig Pep	1	1	1	1	1	1	1	1	1	1	-
AA SEQ ID NO: Y	3930	3931	3932	3933	3934	3935	3936	3937	3938	3939	3940
5' NT of SEQ of First D Start AA of NO: Codon Signal Y	178	46	492	73	254	211	162	404	295	195	89
5' NT of Start Codon	178	46	492	73	254	211	162	404	562	195	89
3' NT of Clone Seq.	3186	1608	1218	898	589	781	829	1007	854	1274	1820
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	I	1	1	1	1	1	1	1	1	—	1
	3186	1608	1218	368	685	781	829	1007	854	1274	1343 1820
NT SEQ ID NO: X	1333	1334	1335	1336	1337	1338	1339	1340	1341	1342	1343
Vector	pCMVSport 3.0 1333	pCMVSport 3.0 1334	pCMVSport 3.0 1335	pCMVSport 3.0 1336	pSport1						
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203957 04/26/99						
cDNA Clone ID	HLWAX50	HLWBJ93	HLWBK16	HLWCC11	HLYAH81	НГ.ҮАН92	HLYAJ79	HLYAL28	HLYAR30	HLYAT54	HLYBC81
Gene No.	1323	1324	1325	1326	1327	1328	1329	1330	1331	1332	1333

Last AA of ORF	30	29	38	9	34	34	33	71	20	35	33
First I AA of AP Of Portion C	29		50	-	20	19	33	19	16	28	17
Last AA of Sig S	78		19		16	18	32	18	15	27	16
First AA of Sig Pep		-	1	1	1	1	1	-			1
AA First SEQ AA ID of NO: Sig Y Pep	3941	3942	3943	3944	3945	3946	3947	3948	3949	3950	3951
5' NT of First AA of Signal Pep	06	497	223	324	237	48	245	369	188	22	46
5' NT of of of Start AA of Codon Signal Pep	06	497	223		237	48	245	369	188	22	46
3' NT of Clone Seq.	1984	789	354	1487	820	751	1084	1403	1566	899	313
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	1	-	26		1	1	1	1
Total NT Seq.	1984	789	354	1487	820	751	1084	1403	1566	899	313
SEQ NO NO X	1344	1345	1346	1347	1348	1349	1350	1351	1352 1566	1353	1354
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203957 04/26/99	203959 04/26/99	203957 04/26/99	203957	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HLYBD09	HLYBL67	HLYBM38	HLYBN23	HLYBN71	HLYBS25	HLYBT28	HLYBU15	HLYBY04	HLYCE15	HLYCH04
Gene No.	1334	1335	1336	1337	1338	1339	1340	1341	1342	1343	1344

Last AA of ORF	46	30	29	12	81	520	37	35	47	46
First AA of Secreted Portion	22	21	28		15	23	31	22	28	18
	21	20	27		14	22	30	21	27	17
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	-	1	1	1	1	1	1	1	-
AA SEQ ID NO: Y	3952	3953	3954	3955	3956	3957	3958	3959	3960	3961
5' NT AA FOOT STATE OF STATE OF STATE AA OF NO: Start AA of NO: Start APP OF NO: Pep	296	180	220	233	427	94	124	20	2789 3960	235
5' NT of Start Codon		180	220	233		94	124	20	2789	235
3' NT of Clone Seq.	1082	1316	722	858	1196	2102	1289	898	2928	1141
5' NT 3' NT of of Total Clone Clone NT Seq. Seq. Seq.	1	1	1	-	46	-	1	-	2604	1
Total NT Seq.	1082	1316	722	858	1206	2102	1289	898	2929	1141
NT SEQ D NO:	1355 1082	1356	1357	1358	1359	1360 2102	1361	1362	1363	1364 1141
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	Uni-ZAP XR
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99
cDNA Clone ID	HLYCY48	HLYDE38	HLYDG55	HLYDO73	HLYEA60	HLYEJ14	HLYEJ44	HLYEU51	HLYGV19	HMABK52
Gene No.	1345	1346	1347	1348	1349	1350	1351	1352	1353	1354

rt l						1		<u> </u>	1
Last AA of ORF	19	34	<i>L</i> 9	257	291	23	64	9	23
First AA of Secreted Portion	18		25	26	39	19	20	23	
Last AA of Sig Pep	17		24	25	38	18	19	22	
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	_	1
AA SEQ ID NO: Y	3962	3963	3964	3965	3966	3967	3968	3969	3970
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	324	251	327	125	264	40	136	743	471
5' NT of Start Codon	324	251		125	264	40	136	743	471
3' NT of Clone Seq.	1285	1796	749	1444	1855	2509	1769	1322	971
5' NT 3' NT of of Clone Clone NT Seq. Seq.	_	-	-	91	619	П	37	695	134
Total NT Seq.	1285	1796	770	1444	1892	2509	2101	1322	1111
NT SEQ ID NO:	1365	1366 1796	1367	1368 1444	1369 1892	1370	1371	1372	1373 1111
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	PTA- 181 06/07/99	203918 04/08/99	203979 04/29/99	203957 04/26/99	PTA- 791 09/27/99	203957 04/26/99	203918 04/08/99	203957 04/26/99
cDNA Clone ID	HMACF34	HMACL77	HMACT74	HMADJ14	HMADJ74	HMAEA58	HMAGF01	HMAJS26	HMCED78
Gene No.	1355	1356	1357	1358	1359	1360	1361	1362	1363

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Last AA of ORF	82	31	11	49	26	98	58	74	82	44
First AA of Secreted Portion	19	29	10	17	17	31	9	11	25	21
Last AA of Sig Pep	18	28	6	16	16	30	5	10	24	20
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep	1	-	-	-	-	-	-			1
	3971	3972	3973	3974	3975	3976	3977	3978	3979	3980
5' NT of First AA of Signal Pep	413	688	317	152	356	215	432	377	399	149
5' NT of Start Codon	413	688			356	215		377	399	
	1514	2780	066	1316	1146	545	909	1035	2414	582
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-	654	1	П	1	1	-	1	273	1
Total NT Seq.	1514	2799	066	1316	1146	545	909	1035	2414	582
NT SEQ ID NO:	1374	1375	1376	1377	1378	1379	1380	1381	1382	1383
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 793 09/27/99	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99	203957 04/26/99	203959 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HMCFN86	HMCGJ47	HMCGK88	HMCIH27	HMCIQ20	HMCJC19	HMDAB44	HMDAE88	HMDAG62	HMDAK20
Gene No.	1364	1365	1366	1367	1368	1369	1370	1371	1372	1373

Last AA of ORF	32	28	137	42	38	32	42	36	50	20	
First AA of Secreted Portion (18		18	13	18	27	21	28	23	14	
	17		17	12	17	26	20	27	22	13	
First AA of Sig Pep	1	1	1	-	1	1		1	1	1	1
AA SEQ ID NO: Y	3981	3982	3983	3984	3985	9868	3987	3988	3989	3990	3991
	31	38	6	141	20	99	192	117	121	553	235
5' NT of of of Start AA of Codon Signal Pep	31	38			20	99	192	117	121		235
3' NT of Clone Seq.	1426	712	1702	1720	505	485	2007	1499	1621	1525	641
5' NT 3' NT of Ol Clone Seq. Seq.	1	1	1	-	-		1	1	1	1	1
Total NT Seq.	1426	712	1702	1720	505	485	2007	1499	1626	2397	641
SEQ BD NO:	1384	1385	1386	1387	1388	1389	1390		1392	1393	1394
Vector	Uni-ZAP XR	Uni-ZAP XR	Lambda ZAP II 1386	Lambda ZAP II 1387	Lambda ZAP II 1388	Lambda ZAP II	Lambda ZAP II 1390 2007	Lambda ZAP II 1391	Lambda ZAP II 1392	Lambda ZAP II	Lambda ZAP II 1394
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	2039 <i>57</i> 04/26/99	203957 04/26/99	203957 04/26/99		203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HMDAM08	HMDAM39	HMEAA41	HMECM77	HMEEH21	HMEET36	HMEEZ07	HMEFB15	HMEIH57	HMEIJ21	HMEIX79
Gene No.	1374	1375	1376	1377	1378	1379	1380	1381	1382	1383	1384

Last AA of ORF	15	36	56	38	58	52	23	21	44	15
First AA of Secreted Portion		30	56	20	34	39	19		30	
Last AA of Sig Pep		29	25	19	33	38	18		29	
AA First SEQ AA ID of NO: Sig Y Pep	_	1	1	1	1	T	1	1	1	1
	3992	3993	3994	3995	3668	3997	3668	3999	4000	4001
	533	124	943	245	26	244	26	243	199	319
5' NT of of of Start AA of Codon Signal Pep	533	124	943	245	26		92	243	661	319
3' NT of Clone Seq.	2163	1312	1933	1726	2006	1175	1339	1221	942	2103
5' NT of Clone Seq.	401	1	710	1	-		-	-		39
Total NT Seq.	2163	1312	1966	1726	2006	1175	1402	1221	942	1404 2103
SEQ NO:	1395	1396		1398	1399	1400	1401	1402	1403	1404
Vector	Lambda ZAP II 1395 2163	Lambda ZAP II 1396	Lambda ZAP II 1397	Lambda ZAP II 1398 1726	Lambda ZAP II 1399	Lambda ZAP II 1400 1175	Lambda ZAP II 1401 1402	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203959 04/26/99	203957 04/26/99		PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203959 04/26/99
cDNA Clone ID	HMEJC96	HMEJD36	HMEJK28	HMEKH55	HMEKW44	HMEKW71	HMELW26	HMGBT32	HMHBI09	HMHBI93
Gene No.	1385	1386	1387	1388	1389	1390	1391	1392	1393	1394

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1 7 0	33	52	39	36	35	38	41	24	8	24	16
A A Sec	23	40	19	19	24	23	28			8	
Last AA of Sig Pep	22	39	18	18	23	22	27			7	
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	_	1	1	1	1	1	1	1	
AA SEQ ID NO:	4002	4003	4004	4005	4006	4007	4008	4009	4010	4011	4012
5' NT of First AA of Signal Pep	208	251	75	273	113	740	212	129	236	360	144
5' NT of Start Codon	208	251	75	273	113	740	212	129	236		
3' NT of Clone Seq.	1255	1642	1621	1978	932	3052	1280	3620	9681	1361	643
5' NT 3' NT of of Clone Clone Seq. Seq.	190	58	1	1	-	597	1	1	T	-	1
Total NT Seq.	1255	1642	1621	1978	932	3052	1280	3620	1896	1361	643
NT SEQ ID NO:	1405	1406 1642	1407 1621	1408	1409	1410	1411	1412	1413	1414	1415
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203918 04/08/99	203918 04/08/99	203957 04/26/99	203959 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203959 04/26/99
cDNA Clone ID	HMHBP74	HMIAC52	HMIAD75	HMIAG42	HMIAG55	HMIAG72	HMIAL39	HMIAO82	HMIAR42	HMIAV33	HMIAZ24
Gene No.	1395	1396	1397	1398	1399	1400	1401	1402	1403	1404	1405

st A F ₹F	<u>~</u>	0	8	0	5	34	15	37	30	2
) I	65	8	33	40	35	3		3	3	
First AA of Secreted Portion	28	24	24	14	28	20		21	18	
Last AA of Sig Pep	27	23	23	13	27	19		20	17	
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	—	1	1	1	1	1	1	1	
AA SEQ ID NO: Y	4013	4014	4015	4016	4017	4018	4019	4020	4021	4022
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	983	31	130	276	316	87	455	09	131	269
5' NT of Start Codon		31	130		316	87	455	09	131	269
3' NT of Clone Seq.	1323	2083	917	1014	1720	1730	2018	1020	957	1034
5' NT 3' NT of of Total Clone Clone NT Seq. Seq. Seq.	734	1	1	1		-	296	-	П	1
Total NT Seq.	1323	2083	917	1014	1720	1730	1422 2018	1020	957	1425 1034
SEQ NO:	1416 1323	1417	1418	1419	1420 1720	1421	1422	1423 1020	1424	1425
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	203918 04/08/99	PTA- 792 09/27/99	203957 04/26/99	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HMIBD93	HMIBE95	HMIBG57	HMJAC12	HMKAN71	HMKBA33	HMKCI22	HMKCK32	HMKCP81	HMKCY49
Gene No.	1406	1407	1408	1409	1410	1411	1412	1413	1414	1415

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Last AA of ORF	39	12	35	19	34	34	48	6	35	22
First AA of Secreted Portion	23	12	12		31	31	37		22	19
Last AA of Sig Pep	22	11	11		30	30	36		21	18
AA First Last SEQ AA AA ID of of of NO: Sig Sig Y Pep Pep	1	1	1	-	-		1	1	1	1
AA SEQ ID NO: Y	4023	4024	4025	4026	4027	4028	4029	4030	4031	4032
5' NT of First AA of Signal Pep	290	286	386	142	127	120	276 4029	290	42	234
5' NT of Start Codon	290			142	127	120	276	290	42	234
3' NT of Clone Seq.	1150	1761	616	573	1384	902	419	1025	669	783
5' NT of Clone Seq.	1	1	1	-	1	1	1	1	1	1
Total NT Seq.	1150	1761	616	573	1384	90/	419	1025	1390	783
NT SEQ ID NO:	1426	1427	1428	1429	1430	1431	1432	1433	1434	1435
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	PTA- 795 09/27/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203918 04/08/99	203957 04/26/99
cDNA Clone ID	HMKDD51	HMKDG69	HMKDM80	HMKEG88	HMMAA09	HMMAK92	HIMMAL32 203957 04/26/99	HMMBD19	HMMBF22	HMMBH91
Gene No.	1416	1417	1418	1419	1420	1421	1422	1423	1424	1425

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Last AA of ORF	20	41	37	4	40	20	42	17	9	33	20
First AA of Secreted Portion		21	22	27	23	24	24			18	
		20	21	26	22	23	23			17	
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	-	1	-	-	-	-	-	-		
AA SEQ B NO:	4033	4034	4035	4036	4037	4038	4039	4040	4041	4042	4043
47 403	Рер 86	260	85	271	103	43	308	236	563	48	24
5' NT of Start Codon	98	260	85	271	103	43		236		48	24
	606	992	712	089	1004	1305	813	1694	865	337	1777
S' NT 3' NT of of Clone Clone Seq. Seq.		П	-	27	П	П	-	_	-	_	-
Total NT Seq.	606	99/	712	089	1004	1305	813	1694	865	337	1777
SEQ NO:	1436	1437	1438	1439	1440 1004	1441	1442	1443	1444	1445	1446 1777
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pBluescript	pBluescript	Uni-ZAP XR
ATCC Deposit No.Z and Date	203957	203957 04/26/99									
cDNA Clone ID	HMMBH94	HMMBK55	HMMBQ31	HMMBR63	HMMBS55	HMMBT47	HMMCD35	HMMCD95	HMPAB26	HMPAP48	HMQAI38
Gene No.	1426	1427	1428	1429	1430	1431	1432	1433	1434	1435	1436

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Last AA of ORF	138	28	ς.	29	3	12	62	4	46	34
First AA of Secreted Portion	26	18		22			15		32	17
	25	17		21			14		31	16
First AA of Sig Pep	1	—	1	1	1	1	1	1	1	1
AA SEQ UO: Y	4044	4045	4046	4047	4048	4049	4050	4051	4052	4053
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	243	57	499	239	288	<i>L</i> 6	221	109	159	316
5' NT of Start Codon	243	57	499	239		26	221			316
3' NT of Clone Seq.	1940	1469	1013	1265	2081	643	809	1277	1982	1600
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	125	1	302		-	1		1	-	157
Total NT Seq.	1940	1469	1449 1013	1450 1265	2122	643	809	1277	1455 1982	1456 1600
NT SEQ ID NO: X	1447	1448 1469	1449	1450	1451	1452	1453	1454	1455	1456
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	2039 <i>57</i> 04/26/99	203957 04/26/99						
cDNA Clone ID	HMQAT69	НМОВГ90	HMQBV82	HMQCA75	НМQСВ37	нмосг80	HMQCX41	НМОДМ09	НМОДО07	HMSAP33
Gene No.	1437	1438	1439	1440	1441	1442	1443	1444	1445	1446

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Last AA of ORF	24	35	35	24	28	21	17	10	28	31
First Last AA of AA Secreted of Portion ORF	22	21	25	6	18			∞	20	16
	21	20	24	8	17			7	19	15
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1		1	1	1	1	-	-	1	-
AA SEQ ID NO:	4054	4055	4056	4057	4058	4059	4060	4061	4062	4063
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	225	231	151	233	240	26	257	132	146	336
5' NT of Start Codon	225	231	151			<i>L</i> 6			146	336
3' NT of Clone Seq.	1818	1264	1366	2077	1993	1932	1541	934	1625	2128
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1	1	1	1	П	1	1	1	1	256
Total NT Seq.	1818	1264	1366	2077	1993	1932	1541	934	1625	1466 2128
NT SEQ D NO:	1457	1458	1459	1460	1461	1462	1463 1541	1464	1465 1625	1466
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 791 09/27/99	203979 04/29/99	203957 04/26/99	203918 04/08/99	203957 04/26/99	203979 04/29/99	203957 04/26/99	203957 04/26/99	203918 04/08/99	PTA- 181 06/07/99
cDNA Clone ID	HMSAZ48	HMSBN18	HMSBS25	HMSBU14	HMSBZ10	HMSCB94	HMSCK12	HMSCP63	HMSCV75	HMSCV85
Gene No.	1447	1448	1449	1450	1451	1452	1453	1454	1455	1456

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Last AA of ORF	37	177	93	13	24	27	18	54	78	3	33
First AA of Secreted Portion	30	49	26	L		61		25	18		19
Last AA of Sig Pep	59	48	25	9		18		24	17		18
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	1	1	-
AA SEQ ID NO: Y	4064	4065	4066	4067	4068	4069	4070	4071	4072	4073	4074
5' NT AA 5' NT of SEQ of First ID Start AA of NO: Codon Signal Y Pep	177	157	851	347	102	154	09	181	738	1556 4073	182
5' NT of Start Codon		157	851		102	154	09	181	738		182
3' NT of Clone Seq.	1309	1686	2153	1790	1319	1504	1645	1466	1828	1171	1507
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1	1	-		1		1	1	1	1	1
Total NT Seq.	1309	1686	2153	1790	1319	1504	1645	1466	1828	1476 2746	1507
NT SEQ ID NO:	1467	1468	1469	1470	1471	1472	1473 1645	1474	1475	1476	1477
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR							
ATCC Deposit No.Z and Date	203957 04/26/99	203979 04/29/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HMSCW44	HMSCZ19	HMSDI67	HMSDI79	HMSDR28	HMSFT25	HMSFW52	HMSGT73	HMSGU30	HMSHB42	HMSHB42
Gene No.	1457	1458	1459	1460	1461	1462	1463	1464	1465	1466	1467

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Last AA of ORF	37	38	22	12	53	49	38	21	26	10
First AA of Secreted Portion	18	29	21	7	20	51	23	16		
	17	28	20	9	16	20	22	15		
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	ī	1	П	1	1	_	П	1	1
AA] SEQ ID NO: Y	4075	4076	4077	4078	4079	4080	4081	4082	4083	4084
5' NT of First AA of Signal Pep	84	488	57	271	222	164	141	132	264	88
5' NT of Start Codon	84	488	57		222	164	141	132	264	88
3' NT of Clone Seq.	1597	1294	2284	1395	1229	1166	2375	1330	1470	1725
5' NT of Clone Seq.	_	ı	-	-		-	-	П	П	1
Total NT Seq.	1597	1294	2284	1395	1229	1166	2375	1330	1470	1725
NT SEQ ID NO:	1478	1479	1480	1481	1482	1483	1484 2375	1485	1486 1470	1487
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR 1487 1725
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 791 09/27/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HMSHN72	HMSHT29	HMSHW73	HMSIC48	HMSII36	HMSIT42	HMSJB08	69IfSWH	HMSJM20	HMSJR44
Gene No.	1468	1469	1470	1471	1472	1473	1474	1475	1476	1477

Last AA of ORF	69	30	31	46	34	32	33	26	32	119
		.,	(')	7					•	
First AA of Secreted Portion	34	70	12	21	25	19	56	23	16	24
Last AA of Sig Pep	33	19	11	20	24	18	25	22	15	23
First AA of Sig Pep	-	-	-	-	-	1	-	1	1	1
AA SEQ ID NO: Y	4085	4086	4087	4088	4089	4090	4091	4092	4093	4094
5' NT AA of SEQ of First ID Start AA of NO: Codon Signal Y	147	93	1554	198	200	209	30	1074 4092	201	247
5' NT of Start Codon	147	93		198	200		30	1074	201	247
3' NT of Clone Seq.	903	1773	2218	1119	1955	1528	2069	1522	1751	752
S' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	122	198	1	-	787	-	-
Total NT Seq.	903	1773	2218	1119	1955	1528	2069	1528	1751	752
NT SEQ ID NO:	1488	1489	1490	1491	1492	1493	1494	1495	1496	1497
Vector	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 3.0 1490	203957 pCMVSport 3.0 1491 1119 04/26/99	pCMVSport 3.0 1492	pCMVSport 3.0 1493	pCMVSport 3.0 1494 2069	pCMVSport 3.0 1495	pCMVSport 3.0 1496	pCMVSport 3.0 1497
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99		203957 04/26/99	203957 04/26/99	203957 04/26/99		203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HMSKQ91	HMSKY45	HMTAF92	HMTAT36	HMUAB93	HMUAD65	HMUAT23	HMUBA47	HMUB122	HMUBK53
Gene No.	1478	1479	1480	1481	1482	1483	1484	1485	1486	1487

Last AA of ORF	17	26	68	30	40	36	40	22	65	34	47
First AA of Secreted Portion		32	5	21	7	18	19	19	22	24	23
Last AA of Sig Pep		31	4	20	9	17	18	-18	21	23	22
First AA of Sig	-	1	1	_	1	1	1	-	-	1	1
	4095	4096	4097	4098	4099	4100	4101	4102	4103	4104	4105
	272	120	256	1085	137	385	69	467	253	323	323
5' NT of of of Start AA of Codon Signal Pep	272	120		1085		385	69	467	253	323	323
	629	608	1208	2141	1118	1769	1149	1281	1149	1732	1730
5' NT 3' NT of of Clone Seq.	1	1	T	006		-	-	1	1	П	-
Total NT Seq.	629	608	1208	2141	1118	1769	1149	1281	1506 1149	1869	1867
SEQ NO:	1498	1499	1500	1501	1502	1503	1504 1149	1505	1506	1507	1508
Vector	pCMVSport 3.0 1498	pCMVSport 3.0 1499	pCMVSport 3.0 1500 1208	pCMVSport 3.0 1501	pCMVSport 3.0 1502 1118	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203979 04/29/99	203918 04/08/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203979 04/29/99	203979 04/29/99
cDNA Clone ID	HMUBN24	HMUB015	HMUBX48	HMUBY57	HMUBZ15	HMVAL15	HMVBC84	HMVBD68	HMVCG17	HMVCS92	HMVCS92
Gene No.	1488	1489	1490	1491	1492	1493	1494	1495	1496	1497	1498

AA First Last BEQ AA AA First Last D of of AA of AA NO: Sig Sig Secreted of Y Pep Pep Portion ORF	6 1 2	7 1 32 33 85	8 1 1 2 144	9 1 13	0 1 26	1 1 17 18 40	2 1 13 14 20	3 1 20 21 63	4 1 5 6 6	5 1 18 19 33
5' NT of First AA of Signal Pep) 179 4106	2 332 4107	2 4108	5 285 4109	340 4110	72 4111	0 410 4112	4 194 4113	268 4114	31 4115
3' NT of Clone Seq.	1156 179	1386 332	1689	1696 285	712 340	486 72	653 410	974 194	472	924 31
Total NT Seq.	11156 1	1472 1	1991 1	1994 1	3 712 1	1 486 1	654 181	5 974 1	7 472 1	3 924 1
NT SEQ ID NO: Vector X	pSport1 1509	pSport1 1510	pSport1 1511	pSport1 1512	Uni-ZAP XR 1513	Uni-ZAP XR 1514	Uni-ZAP XR 1515	Uni-ZAP XR 1516	Uni-ZAP XR 1517	Uni-ZAP XR 1518
ATCC Deposit No.Z and Date	PTA- p3 181 06/07/99	PTA- p. 181 06/07/99		203917 p ² 04/08/99	203957 Uni-	203957 Uni- 04/26/99	203957 Uni- 04/26/99	203957 Uni-	203957 Uni-	_
cDNA Clone ID	HMVDB45	HMVDJ71	HMVDT89	HMVDT89	HMWA065	HMWAO82	HMWBD74	HMWBK35	HMWBK86	HMWBL38
Gene No.	1499	1500	1501	1502	1503	1504	1505	1506	1507	1508

Last AA of ORF	13	40	14	207	33	∞	36	29	28	61	31
First 1 AA of Secreted Portion (12	31		2	28		16	17	19	19	17
	11	30		1	27		15	16	18	18	16
First Last AA AA of of Sig Sig Pep Pep	1	1	1	1	1	1	1	1	1	1	1
	4116	4117	4118	4119	4120	4121	4122	4123	4124	4125	4126
	11	78	116	3	92	127	129	213	190	71	126
5' NT of of of Start AA of Codon Signal Pep		78	116		92	127	129	213	061	71	126
	807	893	1857	1417	1837	493	460	1369	1555	1259	1217
5' NT 3' NT of Ol Clone Seq. Seq.	1	1	1	1	-		1	1		-	1
Total NT Seq.	807	893	2037	1417	1837	493	460	1369	1556	1259	1217
SEQ BD NO:	1519	1520	1521	1522	1523 1837	1524	1525	1526 1369	1527 1556	1528	1529 1217
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203957 04/26/99	203979 04/29/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203979 04/29/99
cDNA Clone ID	HMWBM48	HMWCG28	HMWCP85	HMWDG30	1513 HMWDU20	HMWDX57	HMWDZ63	HMWEA77	HIMWEC03	HMWEF46	HMWEK43
Gene No.	1509	1510	1511	1512	1513	1514	1515	1516	1517	1518	1519

5' NT 3' NT of SEQ AA AA First Last of of 5' NT of SEQ AA AA First ID of of AA of Seq. Seq. Start AA of NO: Sig Sig Secreted Codon Signal Y Pep Pep Portion	6 12 1876 152 152 4127 1 1 2	6 12 1876 135 135 4128 1 16 17	3 560 1133 726 726 4129 1 20 21	9 1 1609 224 224 4130 1 16 17	3 1 1603 86 86 5203 1 20 21	9 1 1359 48 4131 1 25 26	0 1 1490 29 4132 1	1 522 17 17 4133 1 31 32	1 930 185 4134 1 10 11	1 580 209 209 4135 1 18 19
NT SEQ ID Total NO: NT X Seq.	1530 1876	1531 1876	1532 1133	1533 1609	2606 1603	1534 1359	1535 1490	1536 522	1537 930	1538 580
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR 1	Uni-ZAP XR	Uni-ZAP XR 1	Uni-ZAP XR 1	Uni-ZAP XR 1
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	PTA- 181 06/07/99	203959 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203979
cDNA Clone ID	HMWEM23	HMWEM23	HMWER46	HMWEU96	HMWEU96	HMWEX02	HMWFB65	HMWFD77	HMWF025	HMWF089
Gene No.	1520	1521	1522	1523	1523	1524	1525	1526	1527	1528

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Last AA of ORF	318	36	109	32	73	34	69	86	38	46	38
First AA of Secreted Portion	42	29	43	22	28	26	23	21	20	39	38
First Last AA AA of of Sig Sig Pep Pep	41	28	42	21	27	25	22	20	19	38	37
First AA of Sig Pep	_	-	1	-			-		-	_	-
AA SEQ ID NO: Y	4136	4137	4138	4139	4140	4141	4142	4143	4144	4145	4146
5' NT AA of SEQ First ID AA of NO: Signal Y Pep	33	175	41	224	164	53	135	686	115	187	116
5' NT of Start Codon	33	175	41	224	164	53	135	686	115	187	116
5' NT 3' NT of Ol Clone Clone Seq. Seq.	1224	1448	1143	1589	831	784	1178	1579	954	1563	1847
S' NT 3' NT of Of Clone Seq. Seq.	1		-	-	-	-	-	920	1	1	1
Total NT Seq.	1224	1448	1143	1589	831	784	1178	1579	954	1563	1847
NT SEQ ID NO:	1539	1540	1541	1542	1543	1544	1545	1546	1547	1548	1549 1847
Vector	Uni-ZAP XR	pSport1									
ATCC Deposit No.Z and Date	203979 04/29/99	203957 04/26/99	203979 04/29/99	203957 04/26/99							
cDNA Clone ID	HMWGM41	HMWG095	HMWGV85	HMWGZ42	HMWHR36	SSMIMMH	HMWIQ26	HMWIU49	HMWJJ62	HMWJJ64	HNAAD76
Gene No.	1529	1530	1531	1532	1533	1534	1535	1536	1537	1538	1539

Last AA of ORF	23	217	35	40	35	34	4	31	46	23
First I AA of Secreted Portion C	19	41	16	32	27	22	28	21	18	22
Last AA of Sig Pep	18	13	15	31	26	21	27	20	17	21
First AA of Sig	1	1	1	-	1	-	-		-	1
AA SEQ ID NO: Y	4147	4148	4149	4150	4151	4152	4153	4154	4155	4156
	107	88	179	78	112	206	247	99	47	186
5' NT of of of Start AA of Codon Signal Pep	107	88	179	78	112			99	47	186
	1391	1249	1980	772	822	1488	1383	748	1694	1567
S' NT 3' NT of Clone Seq.	-	-				-	105	Н	21	1
Total NT Seq.	1391	1272	2008	772	822	1488	1383	748	1694	1572
NT SEQ ID NO:	1550	1551	1552	1553	1554	1555	1556	1557	1558	1559
Vector	pSport1	pSport1	pSport1	pCMVSport 2.0 1553	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR 1556	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR 1559
ATCC Deposit No.Z and Date	203957 04/26/99	203979 04/29/99	203957 04/26/99		203957 04/26/99	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203979 04/29/99
cDNA Clone ID	HNAAE24	HNALD94	HNALE44	HNDAC35	HNEAA04	HNEAH26	HNEAK38	HNEAK65	HNEBX72	HNEBY79
Gene No.	1540	1541	1542	1543	1544	1545	1546	1547	1548	1549

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Last AA of ORF	46	48	37	37	59	24	99	52	477	8	2
First AA of Secreted Portion	20	33	18	18	18		24	45	2		
Last AA of Sig Pep	19	32	17	17	17		23	44	-	:	
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	1	1_	1	1	1		1	-
AA SEQ D NO: Y	4157	4158	4159	4160	4161	4162	4163	794 4164	4165	4166	4167
5' NT of First AA of Signal Pep	262	1120	71	11	649	211	280	794	227	112	32
5' NT of Start Codon	262	1120	71	71	649		280	794	72Z		
3' NT of Clone Seq.	1265	3135	1314	1293	1564	914	2235	1369	2885	2389	1474
5' NT 3' NT of of Clone Seq. Seq.	1	1077	1	1	575	1	131	1	-	1	480
Total NT Seq.	1265	3332	1314	2545	1564	914	1566 2235	1369	1568 2910	2430	1570 1525
NT SEQ ID' NO: X	1560	1991	1562	1563	1564 1564	1565	1566	1567	1568	1569	1570
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	2039 <i>57</i> 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203957 04/26/99	203917 04/08/99	203917 04/08/99	203959 04/26/99
cDNA Clone ID	HNECD52	HNECL75	HNECX90	HNECX90	HNEDA05	HINEDP75	HNEDQ02	HNEDU46	HNFAD50	HNFAD50	HNFAG67
Gene No.	1550	1551	1552	1553	1554	1555	1556	1557	1558	1559	1560

Last AA of ORF		18	629	34	16	2	15	09	47	27
	17		79	Ω.	1	.,		9	4	2
First AA of Secreted Portion	12		11		14		13	17	19	24
	11		10		13		12	16	18	23
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	-		-	1	-
AA SEQ ID NO: Y	4168	4169	4170	4171	4172	4173	4174	4175	4176	4177
5' NT of First AA of Signal Pep	332	526	317	231	72	30	311	473	6	219
5' NT of Start Codon		526	317	.231	72	30	311			219
3' NT of Clone Seq.	2399	1645	2847	2661	994	793	1226	1336	2253	1481
5' NT of Clone Seq.	1	299	1	-	1	1	1	327	-	1
Total NT Seq.	2399	1709	2847	2661	994	793	1482	1578 1336	1579 2253	1580 1481
SEQ BD NO:	1571	1572	1573	1574	1575	1576	1577 1482	1578	1579	1580
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 793 09/27/99	PTA- 181 06/07/99	PTA- 791 09/27/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203959 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HNFCJ77	HNFCO56	HNFCY57	HNFDL89	HNFDT73	HNFDU92	HNFDY09	HNFDY31	HNFEA17	HNFEP55
Gene No.	1561	1562	1563	1564	1565	1566	1567	1568	1569	1570

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Last AA of ORF	51	30	6	31	89	75	41	12	∞	66
First AA of Secreted Portion	18	18		13	11	47	17			27
Last AA of Sig Pep	17	17		12	10	46	16			26
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	1		1	-	-	-		1	
AA SEQ ID NO: Y	4178	4179	4180	4181	4182	4183	4184	4185	4186	4187
	281	162	217	292	219	12	246	119	281	107
5' NT of of of Start AA of Codon Signal Pep	281		217	292		12	246	119		107
3' NT of Clone Seq.	1268	1637	2127	1551	874	942	1124	1170	1150	2612
5' NT of Clone Seq.	1	-	-	-	-	-	11		-	1
Total NT Seq.	1268	1637	2127	1551	874	942	1124	1170	1150	2612
NT SEQ ID NO:	1881	1582	1583	1584	1585	1586	1587	1588	1589	1590
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99	203959 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	2039 <i>57</i> 04/26/99	203957 04/26/99	PTA- 795 09/27/99
cDNA Clone ID	HINFET12	HNFFR59	HNFGC51	HNFGR15	HNFGW37	HNFGW53	HNFHA34	HNFHD58	HNFHV68	HNFIE15
Gene No.	1571	1572	1573	1574	1575	1576	1577	1578	1579	1580

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Last AA of ORF	19	70	42	32	19	26	35	52	38	70
First AA of Secreted Portion			21	18	16	20	23	19	19	15
Last AA of Sig Pep			20	17	15	19	22	18	18	14
First Last AA AA of of of Sig Sig Pep Pep	-				_	-	1	1	-	
AA SEQ ID NO: Y	4188	4189	4190	4191	4192	4193	4194	4195	4196	4197
5' NT AA 5' NT of SEQ of First ID Start AA of NO: Codon Signal Y Pep	150	168	288	110	168	15	6	145	145	7
5' NT of Start Codon			288	110		15	6	145	145	7
S' NT 3' NT of of Clone Seq. Seq.	1485	1566	1638	935	628	410	1409	1300	397	964
5' NT of Clone Seq.	1	-	-			1	1	1	1	1
Total NT Seq.	1485	1566	1638	935	628	410	1409	1300	397	964
NT SEQ ID NO:	1591	1592	1593	1594	1595	1596	1597	1598	1599	1600
Vector	pBluescript	pBluescript	Uni-ZAP XR	Uni-ZAP XR 1600						
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203957 04/26/99	203959 04/26/99	203957 04/26/99						
cDNA Clone ID	HNFIE29	HNFIG49	HNFJE27	HNFJG16	HNGAC71	HNGAK42	HNGAL25	HNGAT83	HNGAX06	HNGBB09
Gene No.	1581	1582	1583	1584	1585	1586	1587	1588	1589	1590

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Last AA of ORF	55	31	9	4	8	74	36	<i>L</i> 9	17	42
First AA of Secreted Portion	21	14		22	26	25	24	45	5	25
Last AA of Sig Pep	20	13		21	25	24	23	4	4	24
First Last AA AA of of Sig Sig Pep Pep	1		-	1	1	1	-	-	1	-
AA SEQ ID NO: Y	4198	4199	4200	4201	4202	4203	4204	4205	4206	4207
5' NT AA 5' NT of SEQ of First ID Start AA of NO: Codon Signal Y Pep	178	22	208	158	08	180	36	164	135	15
	178	22		158	80	180	36	164	-	15
5' NT 3' NT of Olone Clone Seq.	1004	1110	639	1197	421	1112	418	759	1102	961
S' NT 3' NT of Olone Clone Seq.	1			-	П	-	-	-	2	1
Total NT Seq.	1004	1110	639	1197	421	1112	418	759	1440	961
NT SEQ ID NO: X	1601	1602	1603	1604	1605	1606	1607	1608	1609 1440	1610
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 793 09/27/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HNGBC53	HNGBD94	HNGBE44	HNGBE63	HNGBI83	HNGBJ74	HNGBP30	HNGBQ61	HNGBS35	HNGBW25
Gene No.	1591	1592	1593	1594	1595	1596	1597	1598	1599	1600

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Last AA of ORF	35	25	92	83	38	54	22	41	35	71
First AA of Secreted Portion	21	24	24	28	17	28	61	18	20	25
Last AA of Sig Pep	20	23	23	27	16	27	81	17	19	24
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	1	1
AA SEQ ID NO:	4208	4209	4210	4211	4212	4213	4214	4215	4216	4217
5' NT of First AA of Signal Pep	202	239	36	389	65	78	705	23	206	220
	202	239	36		65	78		23	206	220
3' NT of Clone Seq.	1174	1939	731	1373	1000	1122	847	2111	473	1478
5' NT of Clone Seq.	-	П	1	1			423	T	—	
Total NT Seq.	1174	1939	731	1374	1000	1122	966	2111	473	1478
NT SEQ D NO:	1611	1612	1613	1614 1374	1615	1616	1617	1618	1619	1620
Vector	Uni-ZAP XR									
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99
cDNA Clone ID	HNGCF29	HNGCF64	HINGDF54	HNGDH22	HNGDH27	HNGDN07	HNGDO65	HNGDR39	HNGDW78	HNGEA90
Gene No.	1601	1602	1603	1604	1605	1606	1607	1608	1609	1610

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Last AA of ORF	4	35	31	23	29	20	35	38	89	32
First AA of Secreted Portion		30	22	22	23		56	22	18	23
Last AA of Sig Pep		29	21	21	22		25	21	17	22
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	1	П	1	1	-	-	1
AA SEQ ID NO: Y	4218	4219	4220	4221	4222	4223	4224	4225	4226	4227
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	220	29	36	238	99	241	144	57	178	53
5' NT of Start Codon	220	29	36	238	56	241	144	57	178	53
3' NT of Clone Seq.	601	1120	755	2388	1245	431	1011	569	1223	1626
5' NT 3' NT of of Clone Clone Seq.	1	1	1	1	1	1	1	1		1
Total NT Seq.	601	1120	755	2388	1245	431	1011	569	1223	1630 1626
NT SEQ D NO:	1621	1622	1623	1624	1625	1626	1627	1628	1629	1630
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 792 09/27/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99
cDNA Clone ID	HNGEC17	HNGEE06		HNGEF72	HNGEI64	HNGEJ33	HNGEK64	HNGEN32	HNGER85	HNGES90
Gene No.	1611	1612	1613	1614	1615	1616	1617	1618	1619	1620

Last AA of ORF	54	9	6	25	23	64	5	32	36	18	57
First AA of Secreted Portion	29			18		14		21	20		24
Last AA of Sig Pep	28			17		13		20	19		23
First AA of Sig Pep	1	1	1	1	1	1	1	1	1	1	1
	4228	4229	4230	4231	4232	4233	4234	4235	4236	4237	4238
5' NT AA of SEQ of First ID Start AA of NO: Codon Signal Y	225	389	231	298	466	211	110	14	185	194	250
5' NT of Start Codon	225	389	231	298	466		110	14	185	194	
	1347	741	962	943	1120	402	214	220	1811	462	534
5' NT 3' NT of Of Clone Seq. Seq.	1	1	1	115	306	1	1	1		1	1
Total NT Seq.	1347	741	962	943	1635 1120	402	214	570	1811	462	534
SEQ BD NO:	1631	1632	1633	1634	1635	1636	1637	1638	1639 1811	1640	1641
Vector	Uni-ZAP XR										
ATCC Deposit No.Z and Date	203957 04/26/99										
cDNA Clone ID	HINGET33	HNGEX18	HNGEY45	HNGEZ02	HNGEZ90	HNGFA25	HNGFB05	HINGFD30	HNGFD31	HNGFD61	HNGFG04
Gene No.	1621	1622	1623	1624	1625	1626	1627	1628	1629	1630	1631

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Last AA of ORF	43	99	9	49	99	30	<i>L</i> 9	52	22	33
First Last AA of AA Secreted of Portion ORF	22	21		7	42	22	40	37	18	28
Last AA of Sig Pep	21	20		9	41	21	39	36	17	27
AA First SEQ AA ID of NO: Sig Y Pep	1	-	-	-	-	1	-	-	1	1
AA SEQ ID NO: Y	4239	4240	4241	4242	4243	4244	4245	4246	4247	4248
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	189	963	739	34	220	195	39	73	102	24
		963			220	195	39		102	24
3' NT of Clone Seq.	1011	1665	963	1573	1361	1043	1113	495	1099	029
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-	-	693	-	-	-		-		1
Total NT Seq.	1011	1665	963	1573	1361	1043	1113	495	1099	029
NT SEQ UD NO:	1642	1643	1644	1645	1646 1361	1647	1648	1649	1650	1651
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR				
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203918 04/08/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HNGFG74	HNGFH32	HNGFH83	HNGF121	HNGFM31	HNGFN77	HNGFQ18	HNGFR54	HNGFT70	HNGFU70
Gene No.	1632	1633	1634	1635	1636	1637	1638	1639	1640	1641

Last AA of ORF	6	159	23	38	53	21	37	160	14	-
First AA of Secreted Portion	6	24		26	28	19	20	2		
First Last AA AA of of Sig Sig Pep Pep	8	23		25	27	18	19			
First AA of Sig Pep	1	-	1	1	-	1	1		-	
AA SEQ ID NO:	4249	4250	4251	4252	4253	4254	4255	4256	4257	4258
5' NT of of Start AA of Codon Signal Pep	262	83	225	38	392	43	107	-	153	160
		83	225	38	392	43	107		153	
3' NT of Clone Seq.	1360	840	1590	177	790	1270	1100	1473	1291	582
5' NT of Clone Seq.	Т	П	2	-	197	-	-		-	1
Total NT Seq.	1360	840	1590	177	1014	1270	1100	1473	1291	582
SEQ BD NO:	1652	1653	1654 1590	1655	1656	1657 1270	1658	1659 1473	1660	1661
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99	203979 04/29/99	203957 04/26/99	203918 04/08/99	PTA- 181 06/07/99	PTA- 181 06/07/99	203957 04/26/99	203957 04/26/99
cDNA Clone ID	HNGFV39	HNGGF13	HNGGK63	HNGGK65	HNGGL11		HNGGS92	HNGGT10	HNGGT74	HNGHB89
Gene No.	1642	1643	1644	1645	1646	1647	1648	1649	1650	1651

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Last AA of ORF	36	12	41	31	187	85	19	33	91	148
First AA of Secreted Portion	30		29	20	21	46		17	31	26
Last AA of Sig Pep	29		28	19	20	45		16	30	25
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep	1	-	-	_	-	-		-	-	1
AA SEQ ID NO: Y	4259	4260	4261	4262	4263	4264	4265	4266	4267	4268
5' NT of First AA of Signal Pep	248	234	18	54	221	100	227	49	69	151
5' NT of Start Codon	248	234	18	54	221	100	227	49	69	151
3' NT of Clone Seq.	1219	1543	817	829	783	578	1142	2478	1092	846
5' NT 3' NT of of Of Clone Clone NT Seq. Seq.	1	-			-		_		-	142
=	1219	1543	817	829	783	578	1142	2478	1092	846
NT SEQ ID NO:	1662	1663	1664	1665	1666	1667	1668	1669 2478	1670	1671
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	2039 <i>57</i> 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 795 09/27/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 795 09/27/99
cDNA Clone ID	HNGHD07	HNGHK37	HNGHM47	HNGHT01	HNGHT86	HNGIH40	HNGIK07	HNGIM40	HNGIM83	HNGIO93
Gene No.	1652	1653	1654	1655	1656	1657	1658	1659	1660	1661

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Last AA of ORF	53	28	31	30	32	36	34	30	62	37	55
First AA of Secreted Portion	35	26	27	18	23	20	34	20	31	14	27
Last AA of Sig Pep	34	25	26	17	22	19	33	19	30	13	26
First Last AA AA of of Sig Sig Pep Pep	1	1	1		1	1	-	_	-	-	-
AA SEQ ID NO: Y	4269	4270	4271	4272	4273	4274	4275	4276	4277	4278	4279
5' NT of First AA of Signal Pep		14	155	23	101	229	129	116	147	49	251
5' NT of Start Codon	213	14	155	23	101	229	129	116	147		251
3' NT of Clone Seq.	630	2521	1475	1784	1743	1201	1815	925	921	626	548
5' NT 3' NT of of Clone Clone Seq. Seq.	1	-	_	-	-	-		-	П	_	1
Total NT Seq.	630	2521	1475	1784	1743	1201	1815	925	921	979	548
NT SEQ ID NO: X	1672	1673	1674	1675	1676	1677	1678	1679	1680	1681	1682
Vector	Uni-ZAP XR										
ATCC Deposit No.Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	203918 04/08/99						
cDNA Clone ID	HNGIS27	HNGIU16	HNGIX91	HNGJA68	HNGJB57	HNGJE86	HNGJH26	HNGJJ61	HNGJL07	99SfDNH	HNGJU60
Gene No.	1662	1663	1664	1665	1666	1667	1668	1669	1670	1671	1672

Last AA of ORF	27	132		17	54	34	4	_		T .
	2	13	51	<u></u>	Š	,	34	27	41	
First AA of Secreted Portion	25	26	16		19	22	30		25	
Last AA of Sig Pep	24	25	15		18	21	29		24	
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep	1	-	-	-	-	-	-	-	1	-
AA 1 SEQ ID NO: Y	4280	4281	4282	4283	4284	4285	4286	4287	4288	4289
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	313	98	290	403	299	69	72	54	157	344
5' NT of Start Codon	313	98		403	299	69	72	54	157	344
3' NT of Clone Seq.	526	481	825	455	811	989	1132	1062	675	835
	1	1		1	1	-	1	_	1	-
Total NT Seq.	975	481	825	455	811	636	1132	1062	675	835
NT SEQ ID NO: X	1683	1684	1685	1686	1687	1688	1689	1690	1691	1692
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	PTA- 795 09/27/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HNGKB09	HNGKW35	HNGKY94	HNGLD28	HNGOY36	HNHAB38	HNHAC43	HNHAD34	HNHAG83	HNHAH06
Gene No.	1673	1674	1675	1676	1677	1678	6291	1680	1681	1682

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Last AA of ORF	7	37	2	12	30	36	4	37	4	152
First AA of Secreted Portion		21	S		19	24		17	7	28
Last AA of Sig Pep		20	4		18	23		16	9	27
	1	-	-			-				-
AA SEQ ID NO: Y	4290	4291	4292	4293	4294	4295	4296	4297	4298	4299
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	151	162	705	214	144	TTT	66	243	395	214
5' NT of Start Codon	151	162		214	144		66	243		
3' NT of Clone Seq.	298	1445	888	642	826	2099	953	1559	809	1157
5' NT of Clone Seq.	1	-		-	-		-	1	1	1
Total NT Seq.	298	1445	888	642	826	2099	953	1559	809	1157
NT SEQ D NO:	1693	1694	1695	1696	1697	1698	1699	1700	1701	1702
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR				
ATCC Deposit No.Z and Date	203918 04/08/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203959 04/26/99	203918 04/08/99
cDNA Clone ID	HNHAJ65	HNHAL61	HNHAP58	HNHAW34	HNHAW35	HNHAY26	HNHAY86	HNHAZ20	HNHBE21	HNHBE38
Gene No.	1683	1684	1685	1686	1687	1688	1689	1690	1691	1692

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Last AA of ORF	2	47	75	22	20	30	12	39	38	31	30
First AA of Secreted Portion		20	24	6	32	28		21	27	25	30
Last AA of Sig Pep		19	23	∞	31	27		20	26	24	29
First Last AA AA of of Sig Sig Pep Pep				1	1	-	1	1		-	1
	4300	4301	4302	4303	4304	4305	4306	4307	4308	4309	4310
41 401	rep 657	161	374	293	55	183	226	73	115	32	207
5' NT of Start Codon		161	374			183	226	73	115	32	207
	742	303	1162	759	933	655	366	621	1611	1267	1341
5' NT of Clone Seq.	-	-	-	-	-	1		12	1	1	1
Total NT Seq.	742	303	1162	759	933	655	366	621	1611	1267	1341
NT SEQ ID NO: X	1703	1704	1705	1706	1707	1708	1709	1710	1711	1712	1713
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR 1713
ATCC Deposit No.Z and Date	203959	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203959 04/26/99
cDNA Clone ID	HNHBG18	HNHBI65	HNHBM16	HNHCH78	HNHCP14	HNHCQ44	HNHCT22	HNHCT47	HNHCV48	HNHCZ54	HNHDC52
Gene No.	1693	1694	1695	1696	1697	1698	1699	1700	1701	1702	1703

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Last AA of ORF	14	104	6	30	59	32	34	15	33	20	22
First AA of Secreted Portion		70		16	20	26	24		31	32	16
Last AA of Sig Pep		19		15	19	25	23		30	31	15
AA First Last SEQ AA AA ID of of of NO: Sig Sig Y Pep Pep		1	П	-	1		1	1	-	-	-
<u> </u>	4311	4312	4313	4314	4315	4316	4317	4318	4319	4320	4321
	140	272	1175	94	19	117	291	311	332	170	282
5' NT of Start Codon	140	272		94	19	117	291	311	332	170	282
3' NT of Clone Seq.	372	745	1203	722	252	1195	1347	794	1184	450	1375
S' NT 3' NT of of Clone Clone Seq. Seq.	1	-		_	-	-	-		1	-	1
Total NT Seq.	372	745	1203	722	252	1195	1347	794	1184	450	1375
SEQ BO: NO:	1714	1715	1716	1717	1718	1719	1720	1721	1722	1723	1724 1375
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR							
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HNHDD95	HNHDE58	HNHD117	HNHDL37	HNHDM21	HNHDR57	HNHDR96	HNHDU62	HNHDW34	HNHDX28	HNHDZ06
Gene No.	1704	1705	1706	1707	1708	1709	1710	1711	1712	1713	1714

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Last AA of ORF	- -	16	70	34	20	34	39	181	31	30
First AA of Secreted Portion				18	28	33	18	35	20	21
Last AA of Sig Pep		10		17	27	32	17	34	19	20
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	1	1		1	1		-	-	-
AA SEQ ID NO: Y	4322	4323	4324	4325	4326	4327	4328	4329	4330	4331
5' NT of First AA of Signal Pep	282	180	70	7	219	267	88	29	175	184
	282		70	7	219	267	88	29	175	184
3' NT of Clone Seq.	328	649	1521	653	289	548	872	1782	702	917
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1		-	-	-	-		-		-
Total NT Seq.	328	649	1521	653	687	548	872	1782	702	917
NT SEQ ID NO:	1725	1726	1727	1728	1729	1730	1731	1732	1733	1734
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HNHDZ42	HINHEF37	HNHEF49	HINHEF70	HNHEG30	HNHEH38	HNHEL22	HNHEN70	HNHEP21	HNHEP41
Gene No.	1715	1716	1717	1718	1719	1720	1721	1722	1723	1724

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Last AA of ORF	39	56	78	9	39	32	28	33	38	39	62
First AA of Secreted Portion	21	20	24	:	16	24	27	22	23	21	30
Last AA of Sig Pep	20	61	23		15	23	26	21	22	20	29
First AA of Sig Pep	1	1	1	1	1	1	1	-	1	-	-
AA SEQ ID NO: Y	4332	4333	4334	4335	4336	4337	4338	4339	4340	4341	4342
4, 4,	гер 61	208	131	52	28	307	26	22	244	69	287
5' NT of Start Codon	61	208	131	52	28		26	22	244	69	287
5' NT 3' NT of Ol Clone Clone Seq. Seq.	1260	859	1516	508	311	1338	1736	522	591	610	695
5' NT 3' NT of of Clone Clone Seq. Seq.	-	-		-	1	-	-	-		1	1
Total NT Seq.	1260	859	1516	508	311	1338	1736	522	591	610	695
NT SEQ ID NO:	1735	1736	1737	1738	1739	1740	1741	1742	1743	1744	1745
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203979	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HNHES33	HINHET16	HNHEY29	HNHEZ76	HNHFF60	HNHFF81	HNHFJ49	HNHFR42	HNHFX25	HNHGD95	HNHGR82
Gene No.	1725	1726	1727	1728	1729	1730	1731	1732	1733	1734	1735

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Last AA of ORF	33	29	6	37	13	17	10	99	62	39
First AA of Secreted Portion	17	26		25	12			32	22	24
Last AA of Sig Pep	16	_ 25		24	11			31	21	23
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	_	1		1		-	-	-
AA SEQ ID NO: Y	4343	4344	4345	4346	4347	4348	4349	4350	4351	4352
5' NT of First AA of Signal		298	27	104	142	273	127	294	164	138
)	195	298	27	104	142		127	294	164	138
3' NT of Clone Seq.	568	468	1138	868	764	417	817	1653	713	318
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-	1	_		-	1	7	-	П	
Total NT Seq.	568	468	1138	868	764	417	817	1653	713	318
NT SEQ ID NO: X	1746	1747	1748 1138	1749	1750	1751	1752	1753	1754	1755
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203918 04/08/99	PTA- 181	203918 04/08/99	203918 04/08/99	203959 04/26/99	203959	203959	PTA- 181 06/07/99
cDNA Clone ID	HNHGS62	HNHGY77	HNHHA47	HNHHN22	HNHHW53	HNHIB40	HNHKI74	HNHKV56	HNHLD80	HNHLS76
Gene No.	1736	1737	1738	1739	1740	1741	1742	1743	1744	1745

Last AA of		34	63	36	55	33	46	31	99	47	49
First AA of Secreted	Portion	18	22	22	27	50	18		22	24	22
	rep	17	21	21	26	19	17		21	23	21
	rep	-	1		1	1	-	-	-	-	-
AA SEQ ID NO:	Y	4353	4354	4355	4356	4357	4358	4359	4360	4361	4362
	Signal Pep	177	51	220	25	250	255	17	539	384	82
5' NT of Start	Codon Signal Pep	177	51	220	25	250	255	17	539	384	82
		1860	1120	1068	1272	536	393	889	1430	1803	1149
5' NT of Clone Seq.		1	1	1	1	-	1	1	1	-	1
	Seq.	1860	1120	1068	1272	536	393	889	1430	1803	1149
SEQ SEQ SO:	X	1756	1757	1758 1068	1759	1760	1761	1762	1763	1764	1765
;	Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1	pCMVSport 3.0 1765 1149
ATCC Deposit No.Z	and Date	203959 04/26/99	203959 04/26/99	203959	203959	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99
cDNA	Clone ID	HNHLZ47	HNHMP15	HNHMP62	HNHMY76	HNHMZ01	HNHND14	HNHND94	HNHOF09	HNKAA76	HNTAF42
Gene	Š	1746	1747	1748	1749	1750	1751	1752	1753	1754	1755

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Last AA of ORF	358	37	62	91	38	58	16	37	08	76
First AA of Secreted Portion	45	31	42	17	56	32		25	21	
Last AA of Sig Pep	44	30	41	16	28	31		24	20	
First Last AA AA of of Sig Sig Pep	1	1	1	1	1	1	1	1	1	-
AA SEQ ID NO: Y	4363	4364	4365	4366	4367	4368	4369	4370	4371	4372
5' NT of First AA of Signal Pep	178	158	162	132	1255	187	121	09	20	182
5' NT of Start Codon	178	158	162	132	1255	187	121	09	20	182
3' NT of Clone Seq.	2753	1247	1154	4024	2287	755	515	787	1241	1093
5' NT of Clone Seq.	1	1	_	1	231	П	1		1	1
Total NT Seq.	2753	1247	1154	4024	2287	755	522	787	1241	1093
NT SEQ ID NO:	1766	1767	1768	1769 4024	1770 2287	1771	1772	1773	1774	1775 1093
Vector	pCMVSport 3.0 1766	pSport1	pSport1	pSport1	pSport1	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 795 09/27/99	PTA- 793 09/27/99	203918 04/08/99	203959 04/26/99	203959	PTA- 181 06/07/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HNTCG32	HNTNY89	HNTRB25	HNTRQ40	HNTSQ23	HOAAH51	HOAAI76	HOAAJ09	HOAAL10	HOAAU13
Gene No.	1756	1757	1758	1759	1760	1761	1762	1763	1764	1765

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Last AA of ORF	29	71	37	<i>L</i> 9	30	124	9	33	15	42
First Last AA of AA Secreted of Portion ORF	29	29	13	44	28	25		6		17
Last AA of Sig Pep	28	28	12	43	27	24		∞		16
First AA of Sig Pep	1	-	-	1	1		-	-		-
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	4373	4374	4375	4376	4377	4378	4379	4380	4381	4382
5' NT of First AA of Signal Pep		181	15	223	39	201	19	3184 4380	2128	179
5' NT of Start Codon	186	181	15		39	201			2128	179
3' NT of Clone Seq.	553	1503	605	1156	1357	739	991	828	2559	745
5' NT 3' NT of of Otable Clone Clone NT Seq. Seq.	1	-	1	-	П	-	387	-	1919	1
	553	1503	605	1156	1357	739	991	3287	2621	745
NT SEQ ID NO:	1776	1777	1778	1779	1780	1781	1782	1783	1784	1785
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99
cDNA Clone ID	HOABC12	НОАВН36	HOBNA89	HOBNF51	HODAH24	НОДАН46	HODAV25	HODAW64	HODAY17	HODBA45
Gene No.	1766	1767	1768	1769	1770	1771	1772	1773	1774	1775

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Last AA of ORF	34	92	3	13	33	14	2	11	10	36
First AA of Secreted Portion	24	20			13					33
Last AA of Sig Pep	23	61			12					32
AA First SEQ AA ID of NO: Sig Y Pep	1	1	1	1	1	-	-	-	1	1
AA SEQ ID NO: Y	4383	4384	4385	4386	4387	4388	4389	4390	4391	4392
5' NT of SEQ / of First D Start AA of NO: 5 Codon Signal Y F	179	91	89	228	383	241	169	159	130	250
5' NT of Start Codon	179	91	89	228		241	169	159		250
3' NT of Clone Seq.	931	635	1187	921	096	698	799	1804	831	407
5' NT 3' NT of of Otatal Clone Clone NT Seq. Seq.	Ţ	1		-	172	1	1	1	1	1
	931	635	1187	921	096	698	799	1804	831	407
NT SEQ D NO:	1786	1787	1788	1789	1790	1791	1792	1793 1804	1794	1795
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918	203918 04/08/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	PTA- 791 09/27/99	PTA- 795 09/27/99	203959 04/26/99	203979 04/29/99	203959 04/26/99
cDNA Clone ID	HODBC79	НОВВБ79	HODBF12	HODBF86	HODBF91	HODBW34	HODBX93	HODBZ06	HODCA73	HODCV86
Gene No.	9//1	1777	1778	1779	1780	1781	1782	1783	1784	1785

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Last AA of ORF	47	33	57	17	39	35	28	19	72	56	24
First AA of Secreted Portion	20		31		26	26	20	19	23	22	16
Last AA of Sig Pep	19		30		25	25	61	18	77	21	15
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	-	1	1	1	1	1	1	1
AA SEQ ID NO:	4393	4394	4395	4396	4397	4398	4399	4400	4401	4402	4403
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	111	98	186	1540	237	122	834	373	527	389	558
5' NT of Start Codon	111	98	186	1540		122		373	527		
3' NT of Clone Seq.	1255	1768	826	2038	896	1532	1598	1452	1117	1121	2241
5' NT 3' NT of Clone Clone Seq. Seq.	1	1	1	1441	∞		583	1	-	-	439
Total NT Seq.	1255	1768	826	2243	896	1532	1874	1471	1117	1121	1806 2322
NT SEQ ID NO:	1796 1255	1797	1798	1799	1800	1801	1802	1803	1804	1805	1806
Vector	Uni-ZAP XR										
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203917 04/08/99
cDNA Clone ID	HODCY44	HODDB58	HODDG72	HODDJ25	HODDN21	НОДДОЗ1	900ДДОН	HODEA20	НОБЕМЗ8	HODET37	HOEBI94
Gene No.	1786	1787	1788	1789	1790	1791	1792	1793	1794	1795	1796

5' NT 3' NT	Clone of First ID of of AA of Seq. Start AA of NO: Sig Sig Secreted	Codon Signal Y Pep Pep	1330 197 197 4404 1 26 27 44	1100 302 302 4405 1 23	1892 75 75 4406 1 33 34 37	960 403 403 4407 1 18 19 33	1690 501 501 4408 1 19 20 30	385 258 258 4409 1 14 15 17	1634 695 4410 1 6	889 142 142 4411 1 20 21 66	1578 86 86 4412 1 68 69 180	2082 341 341 4413 1 13 14 41	1671 129 129 4414 1 34
5' N.	Total Clone NT Seq.		1330 1	1100 1	1963 56	960 1	1691 450	385 1	1634	889 1	1578 1	2082 1	1671 1
NT	ΑÖ	×	1807	1808	1809	R 1810	1811	R 1812	R 1813 1634	R 1814	2.0 1815	2.0 1816	2.0 1817
		Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 2.0 1815	pCMVSport 2.0 1816	pCMVSport 2.0 1817 1671
ATCC	Deposit No.Z	and Date	203959	203959	203979 04/29/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203979 04/29/99	203918 04/08/99	203979 04/29/99	203918 04/08/99	203918
	cDNA	Clone ID	HOEBJ70	НОЕСВЗЗ	HOECX21	НОЕDE27	НОЕЕК81	НОЕЕZ62	HOEF126	HOEFL74	HOFMF63	HOFMJ65	HOFMK02
	Gene	No.	1797	1798	1799	1800	1801	1802	1803	1804	1805	1806	1807

Last AA of		152	67	46	29	51	114	37	37	206	35
	2	41	18	23	16	16	6	27	28	19	25
	_	40	17	22	15	15	8	26	27	18	24
AA First Last SEQ AA AA D of of NO: Sig Sig	гер		1	1	1	1	1	1	1	1	1
AA SEQ ID NO:	-	4415	4416	4417	4418	4419	4420	4421	4422	4423	4424
5' NT of First AA of	Codon Signal	149	103	159	170	272	30	365	152	040	242
S' NT of Start	COdon	149	103	159	170	272		365	152	640	242
3' NT of Clone Seq.		1142	006	1326	621	2144	1187	1233	1197	5077	1634
5' NT 3' NT of of Clone Clone NT Seq. Seq.		П	1	1	1	1	1	1		1	1
	oed.	1142	006	1326	621	2144	1187	1233	1197	5077	1634
SEQ BO NO:	<	1818	1819	1820	1821	1822	1823	1824	1825	1826	1827
7 (A		pCMVSport 2.0 1818	pCMVSport 2.0 1819	pCMVSport 2.0 1820	pCMVSport 2.0 1821	pCMVSport 2.0 1822 2144	pCMVSport 2.0 1823	pCMVSport 2.0 1824	pCMVSport 2.0 1825	pCMVSport 2.0 1826	pCMVSport 2.0 1827 1634
ATCC Deposit No.Z	and Date	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA	Clone ID	HOFMO16	HOFMP62	HOFMT59	HOFMV22	HOFND06	HOFNY15	HOFNY28	HOFOC41	HOGAA41	HOGAB51
Gene	No.	1808	1809	1810	1811	1812	1813	1814	1815	1816	1817

# A . H								9		
Last AA of ORF	25	37	26	211	19	63	27	216	14	51
First AA of Secreted Portion	15	24	21	25		29	6	21		29
Last AA of Sig Pep	14	23	20	24		28	8	20		28
First Last AA AA of of Sig Sig Pep Pep	-	1	1	11	1	1	1	1	I	1
AA SEQ ID NO: Y	4425	4426	4427	4428	4429	4430	4431	4432	4433	4434
5' NT AA of SEQ First ID AA of NO: Signal Y	1339	225	205	87	713	151	386	167	283	891
5' NT of of of Start AA of Codon Signal Pep	1339	225	205	87	713	151		167	283	891
3' NT of Clone Seq.	1648	1726	1175	1014	1816	734	1392	930	2263	5083
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1058	1	1	П	995		П	-	1	669
Total NT Seq.	1648	1726	1175	1014	1827	734	1392	959	2263	5083
NT SEQ ID NO:		1829	1830			1833	1834	1835	1836	1837
Vector	pCMVSport 2.0 1828	pCMVSport 2.0 1829	pCMVSport 2.0 1830	pCMVSport 2.0 1831	pCMVSport 2.0 1832	203918 pCMVSport 2.0 1833 04/08/99	pCMVSport 2.0 1834	pCMVSport 2.0 1835	pCMVSport 2.0 1836 2263	pCMVSport 2.0 1837
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203979 04/29/99	203959 04/26/99	203979 04/29/99	203959 04/26/99
cDNA Clone ID	HOGAH40	HOGAP06	HOGAR36	HOGAR71	HOGCC26	HOGCD78	HOGCK03	HOGCL01	НОНВВ36	нонвс57
Gene No.	1818	1819	1820	1821	1822	1823	1824	1825	1826	1827

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Last AA of ORF	21	52	38	23	33	31	43	92	4	64
First AA of Secreted Portion		22	25	23	21	28	∞	31	23	5
Last AA of Sig Pep		21	24	22	20	27	7	30	22	4
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	_	-	-	1	1	1	1	-	-	
AA SEQ ID NO:	4435	4436	4437	4438	4439	4440	4441	4442	4443	4444
	338	150	333	126	18	84	295	189	28	25
5' NT of of of Start AA of Codon Signal Pep	338	150	333	126	18	84		189	28	
3' NT of Clone Seq.	1790	829	2574	1579	2202	1556	2185	1643	2665	1233
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.		1	166	1	1	1	1	35	-	12
Total NT Seq.	1790	829	2574	6251	2202	1556	2185	1649	1846 2665	1258
NT SEQ B NO:	1838	1839		1841	1842	1843	1844	1845	1846	1847
Vector	pCMVSport 2.0 1838	pCMVSport 2.0 1839	pCMVSport 2.0 1840	pCMVSport 2.0 1841	pBluescript SK- 1842	pBluescript	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203959 04/26/99	203959 04/26/99	203959	203918 04/08/99	203918	203918 04/08/99	203959 04/26/99	PTA- 791 09/27/99	PTA- 181 06/07/99
cDNA Clone ID	нонвое6	HOHBZ10	НОНСН71	HOHEB48	HONAH67	HOOAC84	HOPBP13	НООВБС21	HORBI80	HORBL77
Gene No.	1828	1829	1830	1831	1832	1833	1834	1835	1836	1837

St A f	T ₆		9	7	2	2		9	2	8
Last AA of ORF	59	20	99	<i>L</i> 9	85	15	∞	26	42	23
First AA of Secreted Portion	26	19	10	46	51			23	13	11
First Last AA AA of of Sig Sig Pep Pep	25	18	6	45	20			22	12	10
First AA of Sig Pep	-	-	-	1	1		-	-	-	-
AA SEQ ID NO: Y	4445	4446	4447	4448	4449	4450	4451	4452	4453	4454
5' NT AA 5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	Pep 173	188	227	20	81	214	134	62	514	158
5' NT of Start Codon	173	188		20	81	214		62		
3' NT of Clone Seq.	1027	1246	1019	1309	2255	1659	2845	1647	640	902
5' NT 3' NT of of of Clone Clone NT Seq. Seq.	1	127	1	Ţ		-	1	1	1	1
Total NT Seq.	1027	1248	1019	1309	2255	1659	2845	1647	640	706
NT SEQ ID NO:	1848	1849	1850	1851	1852	1853	1854 2845	1855	1856	1857
Vector	Uni-ZAP XR	Uni-ZAP XR 1849	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR				
ATCC Deposit No.Z and Date	203918	203959 04/26/99	203959 04/26/99	203979 04/29/99	203918 04/08/99	203918 04/08/99	PTA- 791 09/27/99	203918 04/08/99	203918 04/08/99	203959 04/26/99
cDNA Clone ID	HOSBX14	HOSCZ41	HOSEM81	HOSEO83	HOSFR35	HOUAZ32	HOUBC29	HOUBG39	HOUCD12	HOUDB17
Gene No.	1838	1839	1840	1841	1842	1843	1844	1845	1846	1847

, r-	-	1	T	1	Τ	Т	1	1	1	1
Last AA of ORF	12	32	14	70	53	18	61	28	26	12
First AA of Secreted Portion		25		16	38	14	13	20	21	
Last AA of Sig Pep		24		15	37	13	12	19	70	
First Last AA AA of of Sig Sig Pep Pep	-		1	1		-	1	1	П	-
AA SEQ ID NO: Y	4455	4456	4457	4458	4459	4460	4461	4462	4463	4464
7	360	83	157	295	376	1509 4460	343	114	228	49
5' NT of Start Codon		83	157	295	376	1509	343	114	228	49
3' NT of Clone Seq.	1152	2249	1450	1645	626	2569	1117	098	1082	696
5' NT 3' NT of of Clone Clone NT Seq. Seq.		-	-	-	-	1423	-		-	-
Total NT Seq.	1264	2249	1450	1645	626	2952	1117	098	1086	696
NT SEQ ID NO:	1858	1859	1860	1861	1862	1863	1864	1865	1866	1867
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203959	PTA- 181 06/07/99	203959	PTA- 181 06/07/99	203959 04/26/99	203959	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HOUDX40	HOUEF84	HOUEJ43	HOUGS36	нопнозе	HOUIG68	HOUIG92	HOVAD93	HOVAE10	HOVAE36
Gene No.	1848	1849	1850	1851	1852	1853	1854	1855	1856	1857

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Last AA of ORF	37	28	16	32	29	32	91	40	4	21
First AA of Secreted Portion	18	20		30		21	18	34	40	
Last AA of Sig Pep	17	19		29		20	17	33	39	
First Last AA AA of of Sig Sig Pep Pep	1	1	-	1	П	_	_	1	1	1
	4465	4466	4467	4468	4469	4470	4471	4472	4473	4474
5' NT of First AA of Signal Pep	154	129	220	379	149	172	569	20	126	169
	154	129	220	379	149	172	269	20	126	169
3' NT of Clone Seq.	1206	1623	1370	751	2258	953	932	1812	594	606
5' NT; of Clone Seq.	1	1	1	1	1	1	1	1	1	1
	1206	1623	1370	751	2329	953	932	1812	594	606
NT SEQ ID NO:	1868 1206	1869	1870 1370	1871	1872	1873	1874	1875	1876	1877
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	PTA- 794 09/27/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HOVAE82	HOVAJ68	HOVAW46	HOVBB19	HOVBD31	HOVBE81	HOVBI16	HOVBS68	HOVCC73	HOVCF30
Gene No.	1858	1859	1860	1861	1862	1863	1864	1865	1866	1867

Last AA of ORF	46	36	47	37	31	38	310	25	29	31
First AA of Secreted Portion	19	23	46	16	22	28	21	19	24	23
Last AA of Sig Pep	18	22	45	15	21	27	20	18	23	22
First Last AA AA of of Sig Sig Pep Pep	1	1	1	1	1	. 1	1	1	1	1
AA SEQ ID NO: Y	4475	4476	4477	4478	4479	4480	4481	4482	4483	4484
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	147	128	65	41	74	149	257	1308 4482	149	214
5' NT of Start Codon		128	65		74	149	257	1308	149	
3' NT of Clone Seq.	1463	808	1583	352	453	966	2435	2435	895	1320
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1	1	1	29	-	_	1215	1215	1	-
Total NT Seq.	1463	608	1583	352	453	966	2444	2444	895	1320
NT SEQ ID NO:	1878	1879	1880	1881	1882	1883	1884	1885	1886	1887
Vector	pSport1	pSport1	pSport1	Uni-ZAP XR	pBluescript SK- 1882	pBluescript SK- 1883	pBluescript SK- 1884 2444	pBluescript SK- 1885	pBluescript SK- 1886	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	PTA- 181 06/07/99	PTA- 181 06/07/99	203918 04/08/99	203959 04/26/99
cDNA Clone ID	HOVCJ71	HOVCN53	HOVCO53	HPASF94	HPBCG26	HPBCT11	НРВ DЕ33	HPBDE33	HPBDF31	HPCAG17
Gene No.	1868	1869	1870	1871	1872	1873	1874	1875	1876	1877

	OKF	31	36	31	72	58	49	35	36	34	18
First AA of Secreted	Portion	23	27	25	24	27	24	22	22	23	
Last AA of Sig	Pep	22	26	24	23	26	23	21	21	22	
	Pep	-	-	1	1	1	1	1	1	1	-
	Ĭ	4485	4486	4487	4488	4489	4490	4491	4492	4493	4494
S' NT AA of SEQ First ID AA of NO:	Codon Signal Pep	81	265	251	166	173	25	207	533	189	53
5' NT of Start	Codon		265	251	166	173	25	207	533	189	53
3' NT of Clone Seq.		873	1176	531	1221	1264	806	546	1160	308	236
5' NT 3' NT of of Clone Clone NT Seq. Seq.		1	1	1	1	1	1	1	250	1	1
Total NT	Seq.	1227	1176	531	1221	1293	806	546	1160	308	236
NT SEQ D NO:	×	1888	1889	1890	1891	1892	1893	1894	1895	1896	1897
	Vector	Uni-ZAP XR	Uni-ZAP XR	pBluescript SK- 1890	pBluescript SK-	pBluescript SK- 1892 1293	Uni-ZAP XR				
ATCC Deposit No.Z	and Date	203959 04/26/99	PTA- 791 09/27/99	203918 04/08/99	203918 04/08/99		203918 04/08/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203918 04/08/99
cDNA	Clone ID	HPCAG17	HPCAM02	HPDDQ17	НРDDQ28	HPDDT14	HPEAA65	HPEAG24	HPEBA84	HPEBF91	HPEBI09
Gene	No.	1878	1879	1880	1881	1882	1883	1884	1885	1886	1887

Last AA of ORF	10	31	6	3	26	9	25	<i>L</i> 9	38	34
First AA of Secreted Portion		27			19		17	17	26	30
		26			18		16	16	25	29
First Last AA AA of of Sig Sig Pep	1	1	1	1	1	1	1	1	1	1
AA SEQ ID NO: Y	4495	4496	4497	4498	4499	4500	4501	4502	4503	4504
5' NT of First AA of Signal Pep	1	78	973	391	174	140	1088 4501	286	74	149
5' NT of of of Start AA of Codon Signal Pep		78		391	174	140	1088		74	149
3' NT of Clone Seq.	859	1345	1376	1485	486	2401	2966	2184	3852	2604
5' NT 3' NT of of Otatal Clone Clone NT Seq. Seq.	1	1	741	1	1	1	785	1	1	2
	2024	1345	1900 1376	1485	486	2401	1904 2970	2184	1906 3852	2604
NT SEQ ID NO:	1898	1899	1900	1901	1902	1903	1904	1905	1906	1907
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR 1907 2604						
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203979 04/29/99	PTA- 791 09/27/99	203918 04/08/99
cDNA Clone ID	HPFCJ75	HPFCP75	HPFDB66	HPFDD28	HPFDI47	HPIAF35	HPIAK27	HPIAL55	HPIAT18	HPIAZ52
Gene No.	1888	1889	1890	1891	1892	1893	1894	1895	1896	1897

			,							
Last AA of ORF	61	24	<i>L</i> 8	31	12	28	18	18	23	29
First AA of Secreted Portion	38	23	98	22			81	18	23	27
Last AA of Sig Pep	37	22	35	21			<i>L</i> 1	17	22	26
First AA of Sig Pep	1	1	1	1	-	1	1	1	1	1
AA SEQ ID NO: Y	4505	4506	4507	4508	4509	4510	4511	4512	4513	4514
5' NT of First AA of Signal Pep	23	49	184	340	158	153	275	275	55	100
5' NT of Start Codon	23	49	184	340	158	153	275	275	55	100
3' NT of Clone Seq.	3033	2002	1417	1146	1465	1792	1953	1953	4161	1211
5' NT 3' NT of Ol Clone Clone Seq. Seq.	T	П	Ţ	1	П		02	70	1	1
Total NT Seq.	3033	2003	1417	1911 1146	1465	1817	1953	1956	4161	1211
NT SEQ ID C	1908 3033	1909 2003	1910 1417	1161	1912	1913	1914 1953	1915	1916 4161	1917 1211
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 793 09/27/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203979 04/29/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HPIBA07	HPIBA24	HPIBI40	HPIBT19	HPJAA82	HPJAB75	HPJAN76	HPJAN76	HPJAU94	HPJAW78
Gene No.	1898	1899	1900	1901	1902	1903	1904	1905	1906	1907

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Last AA of ORF	112	34	36	34	45	65	37	43	11	32
First AA of Secreted Portion	21	18	18	23	21	16	29	20		16
Last AA of Sig Pep	20	17	17	22	20	15	28	61		15
First Last AA AA of of Sig Sig Pep Pep	1	1				-	-	-	-	1
AA SEQ ID NO: Y	4515	4516	4517	4518	4519	4520	4521	4522	4523	4524
5' NT of First AA of Signal Pep	31	288	195	34	527	1830	199	142	216	147
5' NT of Start Codon	31		195	34	527	1830	199	142	216	147
3' NT of Clone Seq.	1703	3121	1501	2203	2971	5065	1561	270	1045	838
5' NT of Clone Seq.	1	1	1	—		1643	-	_		-
Total NT Seq.	1703	3121	1501	2203	2971	5065	1592	270	1045	838
NT SEQ ID NO: X	1918 1703	1919	1920 1501	1921	1922 2971	1923	1924	1925	1926	1927
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203918	203918 04/08/99	PTA- 793 09/27/99	203918 04/08/99	PTA- 181 06/07/99	203957 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HPJBS16	HPJBU04	HPJCN83	HPJCP75	HPJCV35	HPJCX13	HPLAW13	HPMAI31	HPMBI91	HPMBT05
Gene No.	1908	1906	1910	1911	1912	1913	1914	1915	1916	1917

Last AA of ORF	3	25	10	49	54	39	31	31	38	40	18
First AA of Secreted Portion		20		17	44	23	18	20	31	56	
Last AA of Sig Pep		19		16	43	22	17	19	30	25	
First Last AA AA of of Sig Sig Pep Pep		1	1	1	1	1	1	1	1	1	1
AA SEQ ID NO: Y	4525	4526	4527	4528	4529	4530	4531	4532	4533	4534	4535
5' NT of First AA of Signal Pep	523	169	772	166	327	58	233	172	749	166	95
5' NT of Start Codon	523		772	166	327	58		172	749	991	95
3' NT of Clone Seq.	1355	915	2509	921	1720	2310	1541	1727	2288	1156	2215
5' NT 3' NT of of Clone Clone Seq. Seq.	418	1	989	1	230	1	1	91	649	-	1
Total NT Seq.	1367	915	2509	921	1932 1723	2310	1541	2074	1936 2288	1156	1938 2488
NT SEQ ID NO:	1928	1929	1930 2509	1931	1932	1933	1934 1541	1935	1936	1937	1938
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR				
ATCC Deposit No.Z and Date	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203979 04/29/99
cDNA Clone ID	HPMBW95	HPMCW10	HPMCZ18	HPMDA80	HPMDD27	HPMDF45	HPMDP57	HPMEG72	HPMFM70	HPMFP48	HPMFW01
Gene No.	1918	1919	1920	1921	1922	1923	1924	1925	1926	1927	1928

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Last AA of ORF	88	12	55	08	24	31	39	40	44	2
First AA of Secreted Portion	30		21	17	17	26	36	14	21	
	29		20	16	16	25	35	13	20	
First Last AA AA of of Sig Sig Pep Pep	-		1			1	1	1	1	1
AA SEQ ID NO: Y	4536	4537	4538	4539	4540	4541	4542	4543	4544	4545
5' NT of First AA of Signal Pep	67	12	110	1351	322	57	85	233	164	284
5' NT of of of First Start AA of Codon Signal Pep	67		110	1351	322	57	85		164	284
3' NT of Clone Seq.	1640	1831	2065	2217	1697	1575	549	1397	1228	983
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	99	1229	47	1	1	3	1	10
Total NT Seq.	1640	9661	2067	2226	1758	1575	549	1397	1285	985
NT SEQ ID NO:	1939	1940	1941	1942	1943	1944	1945	1946	1947	1948
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript	pBluescript	Lambda ZAP II	Lambda ZAP II 1946	Lambda ZAP II 1947	Lambda ZAP II 1948
ATCC Deposit No.Z and Date	203918 04/08/99	203959 04/26/99	203917 04/08/99	PTA- 1838 05/09/00	203979 04/29/99	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203959	203959 04/26/99
cDNA Clone ID	HPMGM06	HPMGW43	HPMGY89	HPMKB09	HPMSH26	HPMSH96	HPQAJ25	HPQAJ27	HPQAN50	НРQАО80
Gene No.	1929	1930	1931	1932	1933	1934	1935	1936	1937	1938

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Last AA of ORF	31	31	36	36	4	12	28	74	36	73	6
First AA of Secreted Portion	20	21	20	20			23	31	32	49	
Last AA of Sig Pep	19	20	61	19			22	30	34	48	
First AA of Sig Pep	1	1	1	1	1	1	1	1	1	-	1
AA SEQ ID NO: Y	4546	4547	4548	4549	4550	4551	4552	4553	4554	4555	4556
5' NT of First AA of Signal Pep	118	21	251	251	234	236	212	237	146	66	653
5' NT of Start Codon	118	21	251	251	234	236	212	237	146	66	
3' NT of Clone Seq.	911	1017	822	822	1087	1208	944	1071	263	2930	932
5' NT of Clone Seq.	1		1	1	-	3	8	1	1	1	1
Total NT Seq.	961	1017	822	822	1087	1220	951	1071	563	2930	932
SEQ NO:	1949	1950	1951	1952	1953	1954	1955	1956	1957	1958	1959
Vector	Lambda ZAP II 1949	Lambda ZAP II 1950	Lambda ZAP II 1951	Lambda ZAP II 1952	Lambda ZAP II 1953 1087	Lambda ZAP II	Lambda ZAP II 1955	Lambda ZAP II 1956	Lambda ZAP II 1957	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203918 04/08/99
cDNA Clone ID	HPQAW27	нРQВС90	HPQBJ48	HPQBJ48	нРQВL67	HPQBT17	нРQСF94	HPQCI62	HPQRS74	HPRAD30	HPRCC91
Gene No.	1939	1940	1941	1942	1943	1944	1945	1946	1947	1948	1949

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Last AA of ORF	34	51	30	108	10	11	26	187	71	37
First AA of Secreted Portion	29	32		25				22	19	33
Last AA of Sig Pep	28	31		24				21	18	32
First AA of Sig Pep	-	-		_	-	-	-		-	1
AA SEQ ID NO: Y	4557	4558	4559	4560	4561	4562	4563	4564	4565	4566
5' NT of of of Start AA of Codon Signal	2217	151	171	281	142	135	25	102	266	149
5' NT of Start Codon	2217		171		142		25			149
3' NT of Clone Seq.	2882	1959	1139	1572	772	1481	1377	1173	1098	692
5' NT 3' NT of of Clone Clone Seq. Seq.	2001	1	П	26	П	-	-	_		1
Total NT Seq.	2904	1959	1139	2455	772	1481	1377	1173	1098	692
NT SEQ ID NO:	1960	1961	1962	1963	1964	1965	1966	1967	1968	1969
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript	pBluescript	pBluescript	pBluescript
ATCC Deposit No.Z and Date	203959	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	PTA- 792 09/27/99	203959 04/26/99	203959 04/26/99
cDNA Clone ID	HPRCF40	HPRCF50	HPRCL58	HPRCM72	HPRCS59	HPRCT73	HPRTH56	HPTRE80	HPTR142	HPTRL95
Gene No.	1950	1951	1952	1953	1954	1955	1956	1957	1958	1959

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Last AA of ORF	33	70	38	63	35	44	65	9	24	36
First AA of Secreted Portion	14		14	27	20	21	21		22	17
First Last AA AA of of Sig Sig Pep Pep	13		13	26	19	20	20		21	16
First AA of Sig Pep	1		-	1	-			1	1	-
AA SEQ ID NO: Y	4567	4568	4569	4570	4571	4572	4573	4574	4575	4576
5' NT of First AA of Signal Pep	224	80	210	222	992	144	273	271	74	48
5' NT of Start Codon	224	08	210	222	992			271	74	48
3' NT of Clone Seq.	829	1134	1175	553	1453	475	618	520	1068	906
5' NT 3' NT of of Total Clone Clone NT Seq. Seq.				_	576	-	19	-	40	1
Total NT Seq.	859	1134	1175	553	1463	475	636	520	1506	906
NT SEQ ID NO:	1970	1971	1972	1973	1974 1463	1975	1976	1977	1978	1979
Vector	pBluescript	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript	pBluescript	pBluescript	pBluescript	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	PTA- 795 09/27/99	203959 04/26/99	203959 04/26/99	203918 04/08/99
cDNA Clone ID	HPTRQ52	HPTTH35	HPTT165	HPTTT62	HPTVH24	HPTVH59	HPTVI04	HPTVI96	HPVAA15	HPVAB20
Gene No.	1960	1961	1962	1963	1964	1965	1966	1961	1968	1969

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Last AA of ORF	13	39	39	4	34	59	12	24	130	130
First AA of Secreted Portion		26	23	19	15	30		24	30	30
Last AA of Sig Pep		25	22	18	14	29		23	29	29
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1	1				1	-	1	-	1
AA SEQ ID NO:	4577	4578	4579	4580	4581	4582	4583	4584	4585	4586
5' NT of First AA of Signal Pep	311	91	144	46	305	83	401	83	283	283
	311	91	144			83	401	83	283	283
3' NT of Clone Seq.	774	1236	2071	1467	1201	617	637	1609	1994	1994
5' NT 3' NT of of Otal Clone Clone NT Seq. Seq.	-			-	-	-	262	-	94	94
-	774	1236	1982 2071	1467	1201	617	637	1610	2008	1989 2008
NT SEQ ID NO:	1980	1861	1982	1983	1984	1985	1986	1987	1988	1989
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203979 04/29/99	203979 04/29/99
cDNA Clone ID	HPVAB63	HPVAF86	HPWAH55	HPWAO89	HPWAS27	HPWAT86	HPWAV82	HPWBA36	HPWTF23	HPWTF23
Gene No.	1970	1971	1972	1973	1974	1975	9261	1977	1978	1979

	1	1	ı	ı	1	ı	ı	1	1		
Last AA of ORF	6	30	36	25	59	49	31	36	39	4	35
First AA of Secreted Portion	15	17	23	19	18	34	29	34	22		24
	14	16	22	18	17	33	28	33	21		23
First Last AA AA of of Sig Sig Pep Pep	1	-				-	-	1			1
AA SEQ ID NO:	4587	4588	4589	4590	4591	4592	4593	4594	4595	4596	4597
5' NT of of of Start AA of Codon Signal Pen		73	42	∞	274	103	54	95	3201	175	224
5' NT of Start Codon		73	42	∞		103	54	95	3201	175	224
3' NT of Clone Seq.	2190	240	989	1961	2647	1520	594	933	4561	1142	1317
S' NT 3' NT of of Clone Clone Seq. Seq.	227	-	-						3116	-	1
Total NT Seq.	2190	240	989	1961	2647	1520	594	933	4561	1142	1317
SEQ NO.	1990	1991	1992		1994		9661	1997	1998		2000
Vector	Uni-ZAP XR	pBluescript	pBluescript	pCMVSport 3.0 1993	pCMVSport 3.0 1994	pCMVSport 3.0 1995	pCMVSport 3.0 1996	pCMVSport 3.0 1997	pCMVSport 3.0 1998	pCMVSport 3.0 1999	pCMVSport 3.0 2000 1317
ATCC Deposit No.Z and Date	203979 04/29/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203979 04/29/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203959 04/26/99
cDNA Clone ID	HPWTF53	HPXAB56	HPZAB75	HRAAB26	HRAAC36	HRAAF59	HRAAG89	HRAAO40	HRAAZ12	HRABA19	HRABP28
Gene No.	1980	1981	1982	1983	1984	1985	1986	1987	1988	1989	1990

Last AA of ORF	51	48	21	30	46	24	45	45	38	17	44
First AA of Secreted Portion	30	18		18	35	11	18	18	61		19
Last AA of Sig Pep	29	17		17	34	16	11	17	18		18
First AA of Sig Pep	1	1	1	1	1	-	1	1	1		1
	4598	4599	4600	4601	4602	4603	4604	4605	4606	4607	4608
5' NT of First AA of Signal Pep	268	207	150	124	304	147	<i>L</i> 8	98	248	281	235
5' NT of Start Codon	268	207	150	124	304				248	281	235
	420	1506	1424	1348	849	1519	1292	1292	935	2180	948
S' NT 3' NT of Clone Clone Seq.	1	1	1	1	-	1	1	1	09		-
Total NT Seq.	420	1506	1424	1348	849	1519	1292	1292	935	2180	948
NT SEQ ID NO: X	2001	2002	2003	2004	2005	2006	2007	2008	2009	2010	2011
Vector	pCMVSport 3.0 2001	pCMVSport 3.0 2002	pCMVSport 3.0 2003	pCMVSport 3.0 2004	pCMVSport 3.0 2005	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203917 04/08/99	203959	203959 04/26/99
cDNA Clone ID	HRABU56	HRABZ80	HRACB01	HRACI39	HRADU15	HRDAH04	HRDBA20	HRDBD32	HRDBL01	HRDDM85	HRDDS22
Gene No.	1991	1992	1993	1994	1995	1996	1997	1998	1999	2000	2001

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Last AA of	OKF	27	19	41	34	26	137	106	21	37	25
First AA of Secreted	Portion	15	16	30	28	16	16	17		22	12
Last AA of Sig		14	15	29	27	15	15	16		21	11
	Pep	1	1	1	1	1	1	1	1	-	1
	×	4609	4610	4611	4612	4613	4614	4615	4616	4617	4618
	→	203	67	1193	51	75	74	50	158	162	254
S' NT of Start	Codon		67	1193	51	75	74	50	158	162	254
3' NT of Clone Seq.		844	608	1595	953	1320	617	536	451	272	1346
5' NT 3' NT of of Clone Clone NT Seq. Seq.		1	1	821	8	1	1	-	-	-	1
Total NT	Seq.	844	809	1595	953	1320	617	536	451	272	1346
NT SEQ ID NO:	Χ	2012	2013	2014	2015	2016 1320	2017	2018	2019	2020	2021
	1	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript	ZAP Express	ZAP Express	Uni-ZAP XR
ATCC Deposit No.Z	and Date	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	PTA- 181 06/07/99	PTA- 795 09/27/99	203959	203918 04/08/99	203918 04/08/99
cDNA		HRDEJ86	HRDEQ34	HRDFE30	HRDFT83	HRGCA01	HRGCA06	HRGSE38	HRLAT43	HRLME03	HROAN20
Gene	No.	2002	2003	2004	2005	2006	2007	2008	2009	2010	2011

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Last AA of ORF	35	31	31	57	44	14	31	31	42	57	99
First AA of Secreted Portion	21		27	20	29	13	19	19	29	25	33
Last AA of Sig Pep	20		26	19	28	12	18	18	28	24	32
First AA of Sig Pep	1	1	1	1	1	-	-	-	-	1	1
	4619	4620	4621	4622	4623	4624	4625	4626	4627	4628	4629
5' NT of of of First Start AA of Codon Signal Pep	181	152	242	108	395	320	82	266	959	220	193
5' NT of Start Codon	181	152	242	108	395		82	266	626		193
3' NT of Clone Seq.	638	923	1957	516	1157	1084	175	2845	2576	466	1136
s' NT of Clone Seq.	1	1	П	1	1	-	1		891	1	1
Total NT Seq.	638	923	1957	1870	1157	1084	175	2845	2576	466	1136
NT SEQ ID NO:	2022	2023	2024	2025	2026	2027	2028		2030	2031	2032
Vector	Uni-ZAP XR	ZAP Express	pBluescript SK- 2029	pBluescript SK- 2030 2576	pBluescript SK- 2031	pBluescript SK- 2032					
ATCC Deposit No.Z and Date	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203979 04/29/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HROAP64	HROAS35	HROAY16	HROBJ10	HROBW46	HRODG86	HRSAL26	HRTAE88	HRTAP63	HRTAR24	HSAAN03
Gene No.	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	2022

5° NT 3° NT of SEQ AA A First Last Of STORING Clone Clone of First ID of of AA of AA First NT Seq. Seq. Start AA of NO: Sig Sig Secreted Seq. Seq. Start AA of NO: Sig Sig Secreted Pep	1500 1 1500 115 115 4630 1 19	2384 2 2384 246 246 4631 1	947 1 947 110 110 4632 1 29	2187 365 2187 466 466 4633 1 41	937 1 937 211 211 4634 1	419 1 419 132 4635 1 13	4049 1146 3480 1815 4636 1	1377 1 1377 105 4637 1 26	862 1 667 217 217 4638 1 22	1075 1 1075 166 166 4639 1	695 1 695 185 4640 1
NT SEQ ID NO: Vector X	pBluescript SK- 2033	pBluescript SK- 2034	pBluescript SK- 2035	pBluescript SK- 2036 2187	Uni-ZAP XR 2037	Uni-ZAP XR 2038	Uni-ZAP XR 2039 4049	Uni-ZAP XR 2040	Uni-ZAP XR 2041	Uni-ZAP XR 2042	Uni-ZAP XR 2043
ATCC Deposit cDNA No.Z Clone ID and Date	HSAAS05 203959 p	HSAAW13 203918 p	HSABA15 203918 p	HSABG81 203959 p	HSATA50 203918 04/08/99	HSATA61 203959 04/26/99	HSATG66 203917 04/08/99	HSATI91 203918 04/08/99	HSATR50 203959 04/26/99	HSATT82 203918 04/08/99	
Gene No.	2023 F	2024 H	2025 F	2026 F	2027 F	2028 F	2029 F	2030	2031 F	2032 I	2033 F

Last AA of ORF	32	3	38	36	39	16	35	7	22	31
First L AA of A Secreted O Portion O	61		22	21	29		30		18	23
	18		21	20	28		29		17	22
First Last AA AA of of Sig Sig Pep	-	1	1	1		1	1	1	1	-
AA SEQ ID NO: Y	4641	4642	4643	4644	4645	4646	4647	4648	4649	4650
	37	246	62	161	22	169	98	376	179	63
5' NT of of of Start AA of Codon Signal Pep	37	246	62	191	22	169	86		179	63
	721	1029	995	1288	1492	668	2006	1242	1467	851
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	9	1	-		1	1	1	1	1
Total NT Seq.	721	1029	999	1288	1492	668	2006	1242	1467	851
SEQ BD NO:	2044	2045	2046	2047	2048	2049	2050	2051	2052 1467	2053
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	PTA- 181 06/07/99	203959 04/26/99	PTA- 181 06/07/99	203918 04/08/99	203959
cDNA Clone ID	HSATW67	HSATZ02	HSAUA95	HSAUB89	HSAUI53	HSAUV74	HSAUX39	HSAVA58	HSAVE52	HSAVH32
Gene No.	2034	2035	2036	2037	2038	2039	2040	2041	2042	2043

Last AA of ORF	25	22	38	32	3	∞	13	49	59	28
First AA of Secreted Portion	9	20		20				61	22	19
	5	19		16				18	21	18
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	1	-
	4651	4652	4653	4654	4655	4656	4657	4658	4659	4660
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	236	208	727	18	202	100	£ 5 9	121	101	126
		208	727	18		100	653	121		126
3' NT of Clone Seq.	1266	1623	1441	576	645	1134	9851	1703	1114	624
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	-	1	365		48	1	315	1	1	1
	1266	1623	1441	576	5048	1134	1586	1703	1114	624
NT SEQ ID NO:	2054	2055	2056 1441	2057	2058	2059	2060	2061	2062	2063
Vector	Uni-ZAP XR									
ATCC Deposit No.Z and Date	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203960 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203959 04/26/99
cDNA Clone ID	HSAVM49	HSAV011	HSAV017	HSAVQ13	HSAVR85	HSAVY92	HSAVZ05	HSAWB58	HSAWH36	HSAWM20
Gene No.	2044	2045	2046	2047	2048	2049	2050	2051	2052	2053

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Last AA of ORF	9	57	31	69	6	40	20	34	36	35
First AA of Secreted Portion		29	20	32		28		17	20	17
Last AA of Sig Pep		28	19	31		27		16	19	16
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	1	1
AA SEQ ID NO: Y	4661	4662	4663	4664	4665	4666	4667	4668	4669	4670
5' NT of First AA of Signal Pep	223	1363	59	104	98	164	06	260	158	20
5' NT of Start Codon		1363	59	104	98		06	260	158	20
3' NT of Clone Seq.	533	4015	550	812	868	668	484	1391	1125	362
5' NT 3' NT of of Clone NT Seq. Seq.	1	П	Н	-	-	-	_	_	-	1
	533	4015	550	812	868	668	484	1391	1125	366
NT SEQ ID NO:	2064	2065	2066	2067	2068	2069	2070	2071 1391	2072	2073
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	PTA- 791 09/27/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HSAWM74	HSAWX70	HSAXC22	HSAXI10	HSAXI.49	HSAXL82	HSAXN57	HSAX045	HSAXS06	HSAXS22
Gene No.	2054	2055	2056	2057	2058	2059	2060	2061	2062	2063

Last AA of ORF	62	41	23	48	37	14	32	43	35	22
First L AA of A Secreted O Portion O	21 (70 7	16	30 4	25		17	16	23	21
Last AA Fi of A/ Sig Sec Pep Por							16			
	20	19	15	29	24		1	15	22	20
AA First SEQ AA ID of NO: Sig Y Pep	-	1	}		1	1	7 1	2	1	1
• 2	4671	4672	4673	4674	4675	4676	4677	4678	4679	4680
S' NT AA of SEQ First ID AA of NO: Signal Y Pep	151	139	1900	136	173	218	116	246	88	192
5' NT of of of Start AA of Codon Signal Pep	151	139		136	173	218	116	246	88	192
	1066	509	1267	1073	2157	1057	1626	1692	975	1276
5' NT of Clone Seq.	1	117	1559	1	-		-	-	П	П
Total NT Seq.	1066	909	2076 3116	1073	2195	2079 1057	2080 1626	1692	975	2083 1276
NT SEQ ID NO:	2074 1066	2075	2076	2077 1073	2078	2079	2080	2081	2082	2083
tor	AP XR	Uni-ZAP XR	AP XR	Uni-ZAP XR	ipt SK-	VP XR	Uni-ZAP XR	AP XR	Uni-ZAP XR	Uni-ZAP XR
Vector	Uni-ZAP XR	Uni-Z/	Uni-ZAP XR	Uni-Z/	pBluescript SK- 2078	Uni-ZAP XR	Uni-Z	Uni-ZAP XR	Uni-Z	Uni-Z
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203959 04/26/99	203959	203959 ₁	PTA- 181 06/07/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
A7 De N and	20/4	20 4	07/4	07/4	2 4	P P	2 4	20 7	8 4	8 4
A D	L24	082	R62	P90	J47	0618	C55	A26	Y39	72
cDNA Clone ID	HSAYL24	HSAY082	HSAYR62	HSAZP90	HSBAJ47	HSDBI90	HSDDC55	HSDEA26	HSDEY39	hsdff72
Gene No.	2064	2065	2066	2067	2068	2069	2070	2071	2072	2073

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Last AA of ORF	31	24	31	16	35	625	35	12	192	35
First AA of Secreted Portion	24	21	91		18	2	22	L	2	22
Last AA of Sig Pep	23	20	15		17	1	21	9	1	21
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	1		1	1	1	-	1	1	1	1
AA SEQ BOOK	4681	4682	4683	4684	4685	4686	4687	4688	4689	4690
5' NT of First AA of Signal Pep	<i>4L</i>	<i>1</i> 91	217	92	71	765	<i>L</i> 9	940	399	128
5' NT of Start Codon	74	167	217	92	71	765	<i>L</i> 9		399	128
3' NT of Clone Seq.	1212	575	519	1104	865	3231	3229	1124	3291	3290
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	П	П		П	П	П	T	14	1	1
	1212	575	519	1104	865	3244	3229	1545	3304	3303
NT SEQ ID NO:	2084	2085	2086	2087	2088	2089 3244	2090 3229	2091	2092 3304	2093 3303
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99	203917 04/08/99
cDNA Clone ID	HSDFO08	HSDFR10	HSDGB20	HSDGH56	HSDGM01	HSDGM42	HSDGM42	HSDGM42	HSDGM42	HSDGM42
Gene No.	2074	2075	2076	2077	2078	2079	2080	2081	2082	2083

Last AA of ORF	35	62	28	48	30	47	32	116	45	23
First AA of Secreted Portion	22	20	19	28	20	28	23	30	20	15
Last AA of Sig Pep	21	61	18	27	61	27	22	29	19	14
First AA of Sig Pep	1	1	1	1	1	1	1		1	1
AA SEQ ID NO: Y	1691	4692	4693	4694	4695	4696	4697	4698	4699	4700
5' NT of First AA of Signal Pep	128	229	185	194	147	L89	114	74	714	241
5' NT of Start Codon	128	229	185	194	147	289	114			241
3' NT of Clone Seq.	3291	815	1433	1862	1201	1969	1137	1144	1908	1753
5' NT 3' NT of of Clone Clone NT Seq. Seq.		П	-	-	1	439	1	1	245	1
Total NT Seq.	3304	815	1433	1862	1201	1969	1166	1144	1930	1753
SEQ BD NO:	2094	2095	2096	2097	2098	2099	2100	2101	2102	2103
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR						
ATCC Deposit No.Z and Date	203917 04/08/99	203918 04/08/99	203918 04/08/99	203979 04/29/99	203979 04/29/99	203918 04/08/99	203918 04/08/99	203959	203979 04/29/99	203918 04/08/99
cDNA Clone ID	HSDGM42	HSDHD05	HSDIE51	HSDIK31	HSDIV37	HSDJC96	HSDJE77	HSDJF04	HSDJG47	HSDJH72
Gene No.	2084	2085	2086	2087	2088	2089	2090	2091	2092	2093

Last AA of ORF	36	33	496	63	55	96	25	169	13	14
First L AA of // Secreted Portion C			7	43	45	36	25	46		
	21	24	-	42	44	35	24	45		
First AA of Sig Pep	-	-	1	1	-	1	-	1	1	1
AA SEQ ID NO: Y	4701	4702	4703	4704	4705	4706	4707	4708	4709	4710
5' NT AA of SEQ First ID AA of NO: Signal Y Pep	282	102	7	126	199	120	187	1142	330	268
5' NT of of of Start AA of Codon Signal Pep	282	102		126	199	120	187	1142	330	268
3' NT of Clone Seq.	1501	1450	2329	1593	1583	1434	1710	2255	708	1297
5' NT 3' NT of of Of Clone Clone NT Seq. Seq.	-	1	652	1	1	1	1	1029	-	
Total NT Seq.	1501	1450	2329	1593	1583	1434	2110 1710	2279	708	1297
X SEQ NO.	2104 1501	2105	2106 2329	2107	2108	2109	2110	2111	2112	2113 1297
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript	pBluescript	pCMVSport 1
ATCC Deposit No.Z and Date	PTA- 795 09/27/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203959	203918 04/08/99	PTA- 792 09/27/99
cDNA Clone ID	HSDJL07	HSDJR49	HSDJV24	HSDJV40	HSDKA64	HSDKE82	HSDKF96	HSDZO08	96ÖZGSH	HSEBB18
Gene No.	2094	2095	2096	2097	2098	2099	2100	2101	2102	2103

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Last AA of ORF	48	38	72	57	93	36	21	37	26	4
First AA of Secreted Portion	22	19	31	7	26	14		56	22	
Last AA of Sig Pep	21	18	30	6	25	13		25	21	
First AA of Sig Pep	1	1	1	1	1	1	1	_	1	-
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	4711	4712	4713	4714	4715	4716	4717	4718	4719	4720
	170	155	207	42	187	263	46	74	06	196
5' NT of of of Start AA of Codon Signal Pep	170		207			263	46	74	06	
	1434	1358	4416	1285	1528	1225	1913	2192	855	535
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	20	-	464	-	-	1	1	_	1
Total NT Seq.	1434	1501	4416	1287	1544	1225	1913	2192	1385	556
NT SEQ D NO:	2114	2115 1501	2116 4416	2117 1287	2118	2119 1225	2120	2121 2192	2122	2123
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203959 04/26/99	203960 04/26/99	203959 04/26/99	203918 04/08/99	PTA- 181 06/07/99	203959	PTA- 181 06/07/99	PTA- 795 09/27/99	203979 04/29/99
cDNA Clone ID	HSFAM19	HSHAG54	HSHAS72	HSHAX04	HSHBT15	HSHCE85	HSIAC81	HSIAF66	HSIAP01	HSIDA33
Gene No.	2104	2105	2106	2107	2108	2109	2110	2111	2112	2113

Last AA of ORF	57	111	32	15	613	36	14		30	12
	5	11	3	-	[9]	3			3	
First AA of Secreted Portion	20	12	22	:	23	30			18	
Last AA of Sig Pep	61	11	21		22	29			17	
First Last AA AA of of Sig Sig Sig Pep	1	1	1	1	1		1	-		1
AA SEQ ID NO: Y	4721	4722	4723	4724	4725	4726	4727	4728	4729	4730
5' NT AA 5' NT of SEQ of First ID Start AA of NO: Codon Signal Y Pep	260	53	182	162	<u>56</u>	772	259	398	180	189
5° NT of Start Codon			182	162	95	772	259	398	180	
3' NT of Clone Seq.	789	1691	2148	1111	2076	2164	1750	979	2245	1092
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1			П	647	735		-	127	1
Total NT Seq.	789	1691	2148	1111	2150	2238	1750	626	2367	1092
NT SEQ BD NO:	2124	2125	2126	2127	2128	2129	2130 1750	2131	2132	2133 1092
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR						
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203979 04/29/99	203979 04/29/99	PTA- 791 09/27/99	203959 04/26/99	203917 04/08/99	203918 04/08/99
cDNA Clone ID	HSIDA39	HSIDZ25	HSIEB64	HSIEM18	HSIFO61	HSIFO61	HSIGC63	HSIGM95	HSJAE76	HSJAN83
Gene No.	2114	2115	2116	2117	2118	2119	2120	2121	2122	2123

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Last AA of ORF	44	31	181	4	4	4	4	39	20	97
First AA of Secreted Portion	32		36					26		11
Last AA of Sig Pep	31		35				:	25		10
First Last AA AA of of Sig Sig Pep Pep	-	П	-	1	-	1	1	1	-	1
AA SEQ ID NO: Y	4731	4732	4733	4734	4735	4736	4737	4738	4739	4740
5' NT AA I of SEQ of First ID Start AA of NO: Codon Signal Y	99	35	87	416	416	416	416	99	79	235
5' NT of Start Codon	99	35		416	416	416	416	99	79	
3' NT of Clone Seq.	954	541	1142	1349	1349	1349	1349	1195	542	549
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1	-	1	380	380	380	380	-	-	-
Total NT Seq.	954	541	1142	1452	1452	1452	1452	1195	542	549
NT SEQ ID NO: X	2134	2135	2136	2137	2138	2139	2140 1452	2141	2142	2143
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99
cDNA Clone ID	HSJAQ10	HSJAR59	HSJAU93	HSJAY14	HSJAY14	HSJAY14	HSJAY14	HSJBB27	HSKBU03	нѕксо51
Gene No.	2124	2125	2126	2127	2128	2129	2130	2131	2132	2133

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Last AA of ORF	19	17	36	38	81	50	16	46	26	51
First AA of Secreted Portion		14	29	21	21	25		25	17	21
Last AA of Sig Pep		13	28	20	20	24		24	16	20
First Last AA AA of of Sig Sig Pep Pep	-	-	1	1		-	-	-	-	-
AA SEQ ID NO:	4741	4742	4743	4744	4745	4746	4747	4748	4749	4750
5' NT AA of SEQ of First ID Start AA of NO: Codon Signal Y		740	183	315	2066 4745	121	133	1409 4748	66	18
	132		183	315	2066	121	133	1409	66	18
3' NT of Clone Seq.	1707	1159	096	1058	2631	1879	1631	3382	1408	583
5' NT of Clone Seq.	1	474	-	244	1750	-	-	1385		1
Total NT Seq.	1707	1159	096	1065	2631	1879	1631	3382	1408	583
SEQ N	2144 1707	2145	2146	2147 1065	2148	2149	2150 1631	2151	2152	2153
Vector	Uni-ZAP XR	Uni-ZAP XR	pBluescript	pBluescript	pBluescript	pBluescript	pBluescript	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	203959 04/26/99	203979 04/29/99	203959 04/26/99	203959 04/26/99	PTA- 795 09/27/99	PTA- 181 06/07/99	203918 04/08/99	203959 04/26/99	203918 04/08/99
cDNA Clone ID	HSKDE13	HSKDS47	18VHV81	HSKXB14	HSKYR49	HSKYU81	HSKYY92	HSLAB11	HSLAS96	HSLAW59
Gene No.	2134	2135	2136	2137	2138	2139	2140	2141	2142	2143

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Last AA of ORF	61	115	13	345	107	42	69	350	36	58
l sted	25	29		26	50	38	12	27	29	50
	24	28		25	49	37	=	76	28	19
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	-	1	—	1	1	1	-	-	-	
	4751	4752	4753	4754	4755	4756	4757	4758	4759	4760
5' NT of First AA of Signal Pep	58	101	298	13	61	l i	153	270	87	320
5' NT of of of Start AA of Codon Signal Pep		101	598	13	61	596	153	270	87	320
	570	2369	1936	1879	1089	2500	1419	2043	1484	2865
5' NT 3' NT of Clone Clone Seq.	1	1	243		1	746	1	1	1	1
Total NT Seq.	570	2369	1936	1879	1089	2957	1419	2043	1484	2163 2865
SEQ BD NO:	2154	2155	2156	2157	2158	2159	2160	2161	2162	2163
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	PTA- 795 09/27/99	203959	203918 04/08/99	203918 04/08/99	203959	PTA- 181 06/07/99	203959 04/26/99	203918 04/08/99	PTA- 181 06/07/99
cDNA Clone ID	HSLCH54	HSLCH57	HSLCI86	HSLCS31	HSLCS34	HSLCV16	HSLDW54	HSLEC18	HSLEG59	HSLFR59
Gene No.	2144	2145	2146	2147	2148	2149	2150	2151	2152	2153

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Last AA of	5	10	30	26	15	45	53	4	<u> </u>	4
First AA of Secreted Portion		10	17	20	13	24	14			31
Last AA of Sig Pep		6	16	19	12	23	13			30
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	-	-	-	-	1	1		1	1	1
AA SEQ ID NO: Y	4761	4762	4763	4764	4765	4766	4767	4768	4769	4770
5' NT of First AA of Signal Pep	70	226	890	166	262	171	240	226	218	25
5' NT of Start Codon		226	068	166	262	171	240	226		25
3' NT of Clone Seq.	1272	1529	1314	2354	744	372	427	304	400	703
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	834	1	П		1	-	-	-	-	-
Total NT Seq.	1272	1529	1314	2354	744	372	427	304	400	703
SEQ BD NO:	2164	2165	2166	2167 2354	2168	2169	2170	2171	2172	2173
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 792 09/27/99	203959	203959	PTA- 181 06/07/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203959	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HSLGD91	HSLGF66	HSLGF70	HSLGP68	HSNAB88	HSNAH56	HSNAN38	HSNAO19	HSNAQ52	HSNAT08
Gene No.	2154	2155	2156	2157	2158	2159	2160	2161	2162	2163

Last AA of ORF	12	78	34	16	59	14	∞	55	55	12	43
		(4	(4)	_	(4			4,	41	· —	7
First AA of Secreted Portion	6	11	14		28			19	11		16
	∞	10	13		27			18	10		15
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	-	—	1	1	1	1	1	1	-	1	-
AA SEQ ID NO: Y	4771	4772	4773	4774	4775	4776	4777	4778	4779	4780	4781
5' NT of First AA of Signal Pep	174	237	136	286	838	061	231	31	595	131	438
5' NT of Start Codon			136	286	838	190	231	31		131	
3' NT of Clone Seq.	675	432	794	425	1489	323	674	650	757	818	821
5' NT of Clone Seq.	1	1	-	I	781	-		1	1	1	1
Total NT Seq.	912	432	794	425	1489	323	674	650	757	818	821
NT SEQ ID NO:	2174	2175	2176	2177	2178 1489	2179	2180	2181	2182	2183	2184
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203959	203979	203959 04/26/99	203959	203959 04/26/99
cDNA Clone ID	HSNAW06	HSNAW37	HSNBJ05	HSNB090	HSNBQ36	HSNBS39	HSOAE34	HSOAT44	HSOBB94	HSOBH11	HSOBP75
Gene No.	2164	2165	2166	2167	2168	2169	2170	2171	2172	2173	2174

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Last AA of	ORF	18	25	37	37	44	33	46	33	45	13
	Portion		21	22	20	21	20	30	19	22	
Last AA of Sig	Pep		20	21	19	20	19	29	18	21	
First AA of Sig	Pep	1	1	1	-	-	-	-	1	-	1
AA SEQ ID NO:	Y	4782	4783	4784	4785	4786	4787	4788	4789	4790	4791
5' NT of First AA of	Signal Pep	387	252	33	246	261	250	190	179	671	267
5' NT of Start	Codon Signal Pep	387	252	33	246	261	250	190		671	
3' NT of Clone Seq.		735	1372	280	2120	1467	1383	1164	1148	2056	825
5' NT 3' NT of of Clone Clone NT Seq. Seq.		1	1	1	1	-	113	-	-	526	137
Total NT	Seq.	735	1372	580	2120	1467	1917	1164	1180	2056	825
~ ~	×	2185	2186	2187	2188	2189	2190 1917	2191	2192	2193	2194
	Vector	Uni-ZAP XR	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z	and Date	203959 04/26/99	203918 04/08/99	203918 04/08/99	PTA- 793 09/27/99	203918 04/08/99	PTA- 181 06/07/99	203959 04/26/99	203959 04/26/99	PTA- 791 09/27/99	203959 04/26/99
cDNA	Clone ID	HSOBW65	HSPAA89	HSPAC13	HSPAG75	HSPAI20	HSPAL59	HSPAY90	HSPMF63	HSQAC69	HSQAH14
Gene	No.	2175	2176	2177	2178	2179	2180	2181	2182	2183	2184

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Last AA of ORF	81	218	30	12	27	22	12	27	31	30
First AA of Secreted Portion	34	2	26		23	18			20	8
Last AA of Sig Pep	33	-	25		22	17			19	7
First Last AA AA of of Sig Sig Pep Pep	1		1	-	_	1	1	-	-	-
AA SEQ ID NO: Y	4792	4793	4794	4795	4796	4797	4798	4799	4800	4801
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	988	183	41	440	192	302	1001	93	31	267
•	988	183	41	440	192			93	31	
3' NT of Clone Seq.	3107	918	588	2317	1290	1071	1905	1	1469	567
5' NT of Clone Seq.	814	405	-	280	-	188	883	0	7	-
Total NT Seq.	3107	939	588	2317	1290	2290	1934	357	1469	267
NT SEQ ID NO:	2195	2196	2197	2198	2199	2200	2201	2202	2203	2204
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203959 04/26/99	203918 04/08/99	PTA- 793 09/27/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203917 04/08/99	PTA- 181 06/07/99	203918 04/08/99
cDNA Clone ID	HSQAX94	HSQBL20	НЅQСQ45	HSQCY74	HSQDM74	HSQEG23	HSQEG47	HSQFE72	нѕQғе76	HSQFV12
Gene No.	2185	2186	2187	2188	2189	2190	2191	2192	2193	2194

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Last AA of ORF	43	20	14	73	117	34	27	28	14	99	39
First AA of Secreted Portion	19	20		28	21	23		21		20	19
Last AA of Sig Pep	18	19		27	20	22		20		19	18
First Last AA AA of of Sig Sig Pep	1	1	1	1	-	1	1	1	1	_	1
AA SEQ ID NO: Y	4802	4803	4804	4805	4806	4807	4808	4809	4810	4811	4812
5' NT of First AA of Signal Pep	235	131	301	134	8	172	730	<i>L</i> 9	208	308	130
5' NT of Start Codon	235			134		172	730	<i>L</i> 9	208	308	130
	1679	1598	824	2023	942	884	2282	1793	785	844	753
s' NT of Clone Seq.	1	11			-	Π	578				1
Total NT Seq.	1679	1598	824	2023	942	884	2637	1889	785	854	753
NT SEQ ID NO:	2202 1679	2206	2207	2208	2209	2210	2211 2637	2212	2213	2214	2215
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203959	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99
cDNA Clone ID	HSRAA81	HSRAO56	HSRAV28	HSRDM56	HSRDW57	HSREC72	HSREG42	HSRFD18	HSRGZ11	HSRHB59	HSSAN03
Gene No.	2195	2196	2197	2198	2199	2200	2201	2202	2203	2204	2205

St LH LH	<u>,, </u>		7	7	<u>. T</u>	_	350	22	27	71
Last AA of ORF	36	36	92	22	22	27	35	<u>~</u>	2	
First AA of Secreted Portion	16	26	24	17	39	19	25	70	17	33
Last AA of Sig Pep	18	25	23	16	38	18	24	19	16	32
First AA of Sig Pep	-	-	1	1	1			-	1	1
AA SEQ ID NO: Y	4813	4814	4815	4816	4817	4818	4819	4820	4821	4822
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	174	107	261	244	191	166	141	249	261	177
5' NT of Start Codon	174	107	261		191		141	249	261	177
	864	1863	1114	962	1545	1733	1417	1389	1988	1301
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	-	69	1	1	-	-	-	-	
Total NT Seq.	864	1863	1114	962	1545	1733	2222 1417	1389	2224 1988	1301
SEQ NO:	2216	2217	2218 1114	2219	2220	2221 1733	2222	2223	2224	2225
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203959	203959	203918	203959	203918 04/08/99	203918 04/08/99	203979	203918 04/08/99
cDNA Clone ID	HSSCC66	HSSDI13	HSSDQ20	HSSDX38	HSSED57	HSSEL28	HSSFP88	HSSGS62	HSSJA23	HSSJF26
Gene No.	2206	2207	2208	2209	2210	2211	2212	2213	2214	2215

										
Last AA of ORF	57	2	24	24	24	54	13	31	68	=
First AA of Secreted Portion	50		· 91	16	91	20		27	21	
Last AA of Sig Pep	19		15	15	15	19		26	20	
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	-	1	1	1	1	1	1	-	-	-
	4823	4824	4825	4826	4827	4828	4829	4830	4831	4832
5' NT of First AA of Signal Pep	14	151	713	928	<i>1</i> 901	42	127	96	109	295
	14					42	127	96	109	
3' NT of Clone Seq.	2192	1000	1880	2095	2234	831	972	5691	1320	805
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	Ţ	581	962	935	10	1	1	-	1
	2226 2192	1152	2228 1893	2108	2230 2266	831	972	1695	1320	805
NT SEQ ID NO: X	2226	2227	2228	2229	2230	2231	2232	2233	2234	2235
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203979 04/29/99	203979 04/29/99	203979 04/29/99	203979	203959	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99
cDNA Clone ID	HSSJF96	HSSJM47	HSSJW30	HSSJW30	HSSJW30	HSSMY35	HSTAL93	HSTBG23	HSUAF06	HSUBX67
Gene No.	2216	2217	2218	2219	2220	2221	2222	2223	2224	2225

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Last AA of ORF	91	37	40	36	2	31	15	9	4	28	25
First AA of Secreted Portion	24	33	12	33			:		22	24	17
Last AA of Sig Pep	23	32	11	32					21	23	16
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep	1		1	_		-	-	-	-	1	1
AA SEQ ID NO:	4833	4834	4835	4836	4837	4838	4839	4840	4841	4842	4843
5' NT of SEQ of First D Start AA of NO: Codon Signal Y	158	09	295	308	162	309	139	146	120	188	461
5' NT of Start Codon	158	09		308		309	139	146	120		461
3' NT of Clone Seq.	1538	869	525	861	571	446	780	464	785	410	1243
5' NT of Clone Seq.	1	1		-	_	-	-	-	_	-	317
Total NT Seq.	1538	869	525	861	571	446	780	464	785	410	1304
SEQ B B NO:	2236	2237	2238	2239	2240	2241	2242	2243	2244	2245	2246 1304
Vector	pBluescript	Uni-ZAP XR									
ATCC Deposit No.Z and Date	203918 04/08/99	203979 04/29/99	203918 04/08/99	203959 04/26/99							
cDNA Clone ID	HSUSB73	HSVAC05	HSVAE42	HSVAL83	HSVAT36	HSVAV02	HSVBA83	HSVBD37	HSVBN46	HSVBY62	HSVBZ53
Gene No.	2226	2227	2228	2229	2230	2231	2232	2233	2234	2235	2236

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Last AA of ORF	23	18	21	26	ν	92	6	100	16	16	36
First AA of Secreted Portion	:		21			31		14		:	13
Last AA of Sig Pep			20			30		13			12
First Last AA AA of of Sig Sig Pep Pep	1	1	1	1	1	1	1	1	1	1	1
AA SEQ ID NO: Y	4844	4845	4846	4847	4848	4849	4850	4851	4852	4853	4854
5' NT of First AA of Signal Pep	259	257	817	132	217	244	347	345	156	1796	95
5' NT of Start Codon	259	257	817	132	217			345	156		
	1248	914	1122	1041	006	1536	1388	1769	1502	2199	1385
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	-	627	-		94	-	-	-	1413	1.
Total NT Seq.	1248	914	1122	1041	006	1536	1388	1769	1502	2199	1385
NT SEQ ID NO:	2247	2248	2249	2250	2251	2252	2253	2254 1769	2255 1502	2256	2257 1385
Vector	Uni-ZAP XR	pCMVSport 3.0 2248	pCMVSport 3.0 2249	Uni-ZAP XR							
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99
cDNA Clone ID	HSVCF53	HSWAZ17	HSWBI16	HSXAI44	HSXAJ07	HSXAS59	HSXAX20	HSXAY60	HSXBB78	HSXCA83	HSXCX20
Gene No.	2237	2238	2239	2240	2241	2242	2243	2244	2245	2246	2247

Last AA of ORF		34	52	57	47	79	63	168	35	35
First I AA of Secreted Portion C		18	20	24	33	36	22	2	34	14
		17	19	23	32	35	21	1	33	13
First Last AA AA of of Sig Sig Pep Pep	1	1	1	-	1	1	1	1	1	1
AA SEQ ID NO: Y	4855	4856	4857	4858	4859	4860	4861	4862	4863	4864
	686	182	329	425	84	52	2055 4861	21	299	1168 4864
5' NT of of of Start AA of Codon Signal Pep				425	84	52	2055	21	299	1168
	3787	1678	1067	2264	778	3268	3315	1189	1029	2319
5' NT 3' NT of of Total Clone Clone NT Seq. Seq.	738	1	-	9	-	П	1954	95		959
Total NT Seq.	3787	1705	1067	2270	778	3268	2264 3350	3054	1029	2267 2319
SEQ NO:	2258	2259	2260	2261	2562	2263	2264	2265	2266	2267
Vector	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 3.0 2260	pCMVSport 3.0 2261	pCMVSport 3.0 2262	pCMVSport 3.0 2263	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917 04/08/99	203959 04/26/99		PTA- 791 09/27/99	203959 04/26/99	203959 04/26/99	203979 04/29/99	203917 04/08/99	203918 04/08/99	203959 04/26/99
cDNA Clone ID	HSXFG21	HSXFH82	HSYBD33	HSYBR79	HSYBV44	HSYBZ94	HT3AB13	HT4SB02	HT4SB37	HT4SB81
Gene No.	2248	2249	2250	2251	2252	2253	2254	2255	2256	2257

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Last AA of ORF	35	35	52	15	18	215	39	24	21	200	51
First AA of Secreted Portion	14	14	33			18	29	24	17	17	17
	13	13	32			17	28	23	16	16	16
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	-	1	1	-		-	-	
AA SEQ D NO: Y	4865	4866	4867	4868	4869	4870	4871	4872	4873	4874	4875
5' NT of First AA of Signal Pep	1171	1171	86	255	202	188	217	205	287	626	183
5' NT of Start Codon	1171	1171	86	255	202		217	205		626	183
3' NT of Clone Seq.	2320	2320	643	1620	1095	2406	1127	378	2056	2366	2761
5' NT of Clone Seq.	396	796	1	1	1	1	1	1	-	-	1
Total NT Seq.	2331	2331	643	1620	1095	2458	1127	378	2056	2366	2761
NT SEQ ID NO: X	2268	2269	2270	2271	2272	2273	2274 1127	2275	2276	2277	2278 2761
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	203959 04/26/99	203959	203918 04/08/99	203959	203979 04/29/99	203959 04/26/99	203918 04/08/99	203979 04/29/99	203918 04/08/99	203979 04/29/99
cDNA Clone ID	HT4SB81	HT4SB81	HT5FX76	HTABF81	HTACX63	HTADC63	HTADO61	HTADQ22	HTAEC59	HTAED89	HTAEF02
Gene No.	2258	2259	2260	2261	2262	2263	2264	2265	2266	2267	2268

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Last AA of ORF	54	35	54	92	38	37	21	41	53	17
First AA of Secreted Portion	16	34	35	22	25	32	16	34	35	
Last AA of Sig Pep		33	34	21	24	31	15	33	34	
First Last AA AA of of Sig Sig Pep Pep	, 1	1	1			1	1	1	1	1
AA SEQ U SO:	4876	4877	4878	4879	4880	4881	4882	4883	4884	4885
5' NT of First AA of Signal		63	167	1414	<i>L</i> 9	290	180	157	83	949
5' NT of of of First Start AA of Codon Signal		63	167	1414	<i>L</i> 9	290	180	157	83	949
	1601	1514	1079	2796	2200	955	1512	946	1570	2418
5' NT 3' NT of of Of Clone Clone NT Seq. Seq.	147	_	1	1188	-	188		1	1	726
Total NT Sea.	1601	1514	1079	2814	2200	996	1512	946	1570	2288 2418
NT SEQ ID NO:	2279 1601	2280	2281	2282	2283 2200	2284	2285	2286	2287	2288
Vector	æ	Uni-ZAP XR	pSport1	pSport1	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959	203959	203918 04/08/99	PTA- 792 09/27/99	PTA- 181 06/07/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203979 04/29/99
cDNA Clone ID	∞	HTAE035	HTDAF68	HTDAI38	HTEAJ87	HTEAN76	HTEBL56	HTECE87	HTEDF78	HTEDT87
Gene	2269	2270	2271	2272	2273	2274	2275	2276	2277	2278

ist f KF	0	0	3	33	49	70	12	35	99	30
Last AA il of ORF	20	30	33	233	4	2	-	3	9	3
First AA of Secreted Portion	19	22	20	2	21			21	18	24
Last AA of Sig Pep	18	21	19	1	20			20	17	23
First AA of Sig Pep	1	1	1	1	1	1	1	1	1	
AA SEQ ID NO: Y	4886	4887	4888	4889	4890	4891	4892	4893	4894	4895
5' NT of First AA of Signal Pep	59	72	156	1	289	36	271	192	343	38
5' NT of Start Codon	99	72	156		289	36	271	192	343	38
3' NT of Clone Seq.	2220	1721	2248	2151	763	1134	963	1738	2186	1316
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-		91	1	1	1	1	-	-	1
Total NT Seq.	2220	1721	2267	2158	763	1134	963	1876	2202	1316
NT SEQ ID NO:	2289	2290	2291	2292	2293	2294	2295	2296	2297	2298
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	PTA- 181 06/07/99	203959	203959 04/26/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	PTA- 181 06/07/99	203918
cDNA Clone ID	HTEDX05	HTEEC19	нтесн03	HTEGH03	HTEGS48	HTEGY81	HTEHB11	нтенв49	HTEHS91	HTEHV60
Gene No.	2279	2280	2281	2282	2283	2284	2285	2286	2287	2288

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Last AA of ORF	29	38	17	24	9	41	29	11	24	9
First AA of Secreted Portion	13	19			,	21	22		24	19
Last AA of Sig Pep	12	18				20	21		23	18
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	1	1
AA First I SEQ AA DO OF SIG SIG I Y Pep I	4896	4897	4898	4899	4900	4901	4902	4903	4904	4905
5' NT of First AA of Signa Pep	96	141	156	183	207	102	96	257	94	124
5' NT of Start Codon		141	156	183		102	96	257	94	124
3' NT of Clone Seq.	1167	1436	1398	673	1051	743	429	1471	1154	395
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	961	·	1	1	1	1	1
Total NT Seq.	1167	1436	2593	673	1051	743	429	1471	1154	395
NT SEQ D NO:	2299	2300 1436	2301	2302	2303	2304	2305	2306 1471	2307	2308
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203959 04/26/99	203917 04/08/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99
cDNA Clone ID	HTEHW80	HTEID25	HTEIJ23	HTEIM62	HTEIV33	HTEIV65	HTEJC50	HTEJD20	HTEJD61	HTEJF31
Gene No.	2289	2290	2291	2292	2293	2294	2295	2296	2297	2298

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Last AA of ORF	17	37	33	4	31	35	322	32	21	142
First AA of Secreted Portion	16	27	21	19	16	16	34	24	16	47
Last AA of Sig Pep	15	26	20	18	15	15	33	23	15	46
AA First Last SEQ AA AA ID of of of NO: Sig Sig Y Pep Pep	1	1	1	-	1	1	1	1	1	1
AA SEQ ID NO: Y	4906	4907	4908	4909	4910	4911	4912	4913	4914	4915
5' NT AA 5' NT of SEQ of First ID Start AA of NO: Codon Signal Y Pep	298	136	389. 4908	1747	41	457	634	146	389	2365 4915
5' NT of Start Codon	298	136	389	1747	41	457	634	146		2365
3' NT of Clone Seq.	975	1158	754	2908	889	930	1663	288	1719	3274
5' NT 3' NT of of Total Clone Clone NT Seq. Seq.	-	-	-	1602		-		-	-	1996
Total NT Seq.	975	1158	754	2908	889	930	1663	288	1719	2318 3299
NT SEQ ID NO:	2309	2310	2311	2312	2313	2314	2315 1663	2316	2317	2318
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1						
ATCC Deposit No.Z and Date	203959 04/26/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	PTA- 792 09/27/99	203959 04/26/99	203959 04/26/99	PTA- 1838 05/09/00
cDNA Clone ID	HTEJI29	HTEJL16	HTEJP65	HTEJY20	HTEKD35	HTEKP82	HTEKV69	HTEKZ52	HTEQG28	HTFOB75
Gene No.	2299	2300	2301	2302	2303	2304	2305	2306	2307	2308

Last AA of ORF	10	23	23	52	34	42	16	34	73	16	28
First AA of Secreted Portion (14		20	24	22		25	33	13	
		13		19	23	21		24	32	12	
First AA of Sig Pep	1	1	1	1	1	1	1	1	1	П	1
AA SEQ D NO:	4916	4917	4918	4919	4920	4921	4922	4923	4924	4925	4926
	180	311	211	127	26	435	155	601	188 4924	271	55
5' NT of of Of Start AA of Codon Signal Pep	180		211	127	26	435	155	109	188	271	55
	1633	068	1074	631	1104	1803	1610	1228	787	1131	1133
5' NT 3' NT of Clone Clone Seq.	25	7	0/9	1		1	1	1	П		
Total NT Seq.	1633	068	1074	631	1104	1803	1610	1228	787	1131	1133
NT SEQ D NO:	2319	2320	2321	2322	2323	2324	2325	2326	2327	2328	2329
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR							
ATCC Deposit No.Z and Date	203959 04/26/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99
cDNA Clone ID	HTGAA35	HTGAD74	HTGAP05	HTGAQ29	HTGAR21	HTGAS70	HTGAT65	HTGAU17	HTGBF47	HTGBK95	HTGCC01
Gene No.	2309	2310	2311	2312	2313	2314	2315	2316	2317	2318	2319

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Last AA of ORF	15	22	39	4	38	27	20	32	14	20
First AA of Secreted Portion	7	18	32	23	31	21		15		18
Last AA of Sig Pep	9	17	31	22	30	20		14		17
First Last AA AA of of Sig Sig Pep Pep	1	1	1	_	1	1	1	1		-
	4927	4928	4929	4930	4931	4932	4933	4934	4935	4936
5' NT of First AA of Signal Pep	231	271	08	13	73	290	243	128	519	335
5' NT of of of Start AA of Codon Signal Pep	231	271	08	13	73	290		128	519	335
	926	950	1325	2301	2057	1927	793	1943	1479	538
5' NT 3' NT of of Of Clone Clone NT Seq. Seq.	208	1	1				1	1	458	1
Total NT Seq.	796	950	1325	2301	2057	1927	793	1943	1479	538
NT SEQ ID NO:	2330	2331	2332	2333 2301	2334	2335	2336	2337	2338	2339
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203918 04/08/99	PTA- 792 09/27/99	PTA- 181 06/07/99	203959	203959 04/26/99	203918 04/08/99	203918 04/08/99	203959 04/26/99
cDNA Clone ID	HTGCK43	HTGDS43	HTGDS92	HTGEX34	HTGFM31	HTGGM37	HTGGN22	HTHAA41	нтнвс58	HTHBO72
Gene No.	2320	2321	2322	2323	2324	2325	2326	2327	2328	2329

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Last AA of ORF	39	12	37	23	7	72	21	80	67	43
First AA of Secreted Portion	22		36			42	14	44	18	18
Last AA of Sig Pep	21		35			41	13	43	17	17
First AA of Sig Pep	1	1	1	1	1	1	1	1	1	
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	4937	4938	4939	4940	4941	4942	4943	4944	4945	4946
5' NT of First AA of Signal	165	207	31	158	250	105	175	118	247	217
5' NT of Start Codon	165	207	31	158		105	175			217
	1090	2025	986	627	1220	657	1026	1535	1242	633
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	-	1	-	-		П	1	1	1	1
Total NT Seq.	1090	2025	986	627	1220	657	1026	1535	1242	633
NT SEQ ID NO:	2340	2341	2342	2343	2344 1220	2345	2346 1026	2347	2348	2349
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	нтнв029	HTHBT76	HTHBZ91	HTHCA30	HTHCM60	HTHDB20	HTHDF45	HTHDF86	HTHDH18	HTHDP65
Gene No.	2330	2331	2332	2333	2334	2335	2336	2337	2338	2339

F F	2	3	2	7	8	0	4	22	ν.	6
Last AA I of ORF	45	23	92	27	33	190	194	232	. 75	49
First AA of Secreted Portion	35	L	29	24	18	15	19	13	16	22
Last AA of Sig Pep	34	9	28	23	17	14	18	12	15	21
First AA of Sig Pep	1	1	1	1	1	-	-	-	-	1
	4947	4948	4949	4950	4951	4952	4953	4954	4955	4956
5' NT of First AA of Signal Pep	09	119	138	263	163	195	176	26	46	263
	09		138	263	163		176	<i>L</i> 6		263
3' NT of Clone Seq.	422	391	2259	854	971	1229	1235	1124	920	489
5' NT 3' NT of Clone Seq.	1	1	-	-	1	1		1	1	1
Total NT Seq.	422	535	2259	854	971	1229	1260	1124	920	489
NT SEQ UD NO:	2350	2351	2352	2353	2354	2355	2356	2357	2358	2359
Vector	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 2.0 2352	pCMVSport 2.0 2353	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	PTA- 181 06/07/99	203959 04/26/99	203959 04/26/99	203979 04/29/99	203959 04/26/99
cDNA Clone ID	HTHDT25	HTHDV50	HTJMA64	HTJMJ72	HTLAD74	HTLAF81	HTLBF46	HTLBF63	HTLCX82	HTLDD89
Gene No.	2340	2341	2342	2343	2344	2345	2346	2347	2348	2349

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Last AA of ORF	93	176	∞	170	13	13	152	75	35
First AA of Secreted Portion	20	24		19			39	19	16
Last AA of Sig Pep	61	23		18			38	18	15
AA First Last SEQ AA AA D of of of NO: Sig Sig Y Pep Pep	1	-	-	-	1	-	1	1	-
AA SEQ ID NO: Y	4957	4958	4959	4960	4961	4962	4963	4964	4965
5' NT AA 5' NT of SEQ of First ID Start AA of NO: Codon Signal Y Pep	84	66	115	110	1559	224	133	276	187
	84	66	115	110	1559	224	133	276	187
3' NT of Clone Seq.	882	957	1240	928	2533	1192	1507	1129	1003
5' NT 3' NT of of Clone Clone Seq. Seq.	_	_	-	_	1336	-	-	-	-
Total NT Seq.	882	957	1240	928	2569	1192	1507	1129	1003
NT SEQ D NO:	2360	2361	2362	2363	2364 2569	2365	2366	2367	2368
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	PTA- 793 09/27/99	PTA- 1838 05/09/00	203918 04/08/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	PTA- 181 06/07/99	203959 04/26/99
cDNA Clone ID	HTLDN34	HTLDP19	HTLDY30	HTLEJ24	HTLEJ75	HTLEJ75	HTLEP55	HTLEV80	HTLEZ57
Gene No.	2350	2351	2352	2353	2354	2355	2356	2357	2358

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Last AA of ORF	314	174	170	82	303	187	187	187	187	314
First AA of Secreted Portion	33	20	29	20	24	21	21	21	21	20
	32	19	28	19	23	20	70	70	70	19
AA First Last SEQ AA AA D of of NO: Sig Sig Y Pep Pep	_	1	-	-	1	1	1	1	1	1
AA SEQ ID NO: Y	4966	4967	4968	4969	4970	4971	4972	4973	4974	4975
	131	763	24	363	337	1413	1413	1413 4973	1413	10
5' NT of of of Start AA of Codon Signal Pep	131	292	24	363	337	1413	1413	1413	1413	10
3' NT of Clone Seq.	1304	1703	699	1189	1245	2133	2184	2133	2133	1082
5' NT of Clone Seq.	1	539	-	-	-	1059	1059	1059	1059	1
Total NT Seq.	1314	1703	699	1189	1245	2204	2240	2240	2240	1082
NT SEQ ID NO:	2369	2370 1703	2371	2372 1189	2373	2374 2204	2375	2376 2240	2377	2378 1082
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203960 04/26/99	203959 04/26/99	PTA- 795 09/27/99	PTA- 795 09/27/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	PTA- 795 09/27/99
cDNA Clone ID	HTLFA90	HTLGL33	HTLGQ25	HTLGS72	HTLGY50	HTLHN86	HTLHN86	HTLHN86	HTLHN86	HTLIW29
Gene No.	2359	2360	2361	2362	2363	2364	2365	2366	2367	2368

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Last AA of ORF	240	31	23	51	51	51	51	28	15	51
First AA of Secreted Portion	39	19	16	23	23	23	23	10		20
Last AA of Sig Pep	38	18	15	22	22	22	22	6		49
First Last AA AA of of Sig Sig Pep Pep	1	1	1	1	-	1	1	1	1	1
AA SEQ ID NO: Y	4976	4977	4978	4979	4980	4981	4982	4983	4984	4985
	966	166	225	1737	1737	1737	1737	551	138	64
5' NT of of of First Start AA of Codon Signal Pep	966	166	225	1737	1737	1737	1737		138	64
3' NT of Clone Seq.	1873	1989	336	2782	2763	2764	2765	1014	1382	1282
5' NT 3' NT of of Clone Clone Seq. Seq.	<i>L</i> 69	1	1	1511	1511	1511	1511	-	-	1
Total NT Seq.	1913	1989	336	2794	2792	3351	2794	1014	1382	1282
NT SEQ ID NO:	2379	2380	2381	2382	2383	2384	2385	2386	2387	2388
Vector	Uni-ZAP XR	pBluescript SK- 2380	pBluescript SK- 2381	pBluescript SK- 2382	pBluescript SK- 2383	pBluescript SK- 2384 3351	pBluescript SK- 2385	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 1838 05/09/00	PTA- 793 09/27/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99
cDNA Clone ID	HTLJC15	HTNAL14	HTNAL34	HTNBJ15	HTNBJ15	HTNBJ15	HTNBJ15	HTOAC65	HTOAE47	HTOAK03
Gene No.	2369	2370	2371	2372	2373	2374	2375	2376	2377	2378

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Last AA of ORF	34	118	4	36	62	14	16	31	19	9
First AA of Secreted Portion	14	20		28	22			23	15	
	13	19		27	21	1		22	14	
First Last AA AA of of Sig Sig Pep	1	1	1	1	1	1	1	1	1	1
AA SEQ ID NO: Y	4986	4987	4988	4989	4990	4991	4992	4993	4994	4995
5' NT of of Of Start AA of Codon Signal Pep	157	82	1123 4988	06	1525 4990	<i>L</i> 9	205	99	139	325
5' NT of Start Codon	157					<i>L</i> 9		99		
3' NT of Clone Seq.	1637	1522	1344	1399	3251	1594	1455	2020	1774	1619
5' NT of Clone Seq.	. 1	1	-	1	1346	1	1	-	-	1
Total NT Seq.	1637	1522	1344	1399	3261	1594	1455	2020	1774	1619
NT SEQ ID NO: X	2389	2390	2391	2392	2393	2394 1594	2395	2396 2020	2397	2398 1619
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HTOAO58	HTOAT56	HTOBG07	HTOBG62	HTODA92	HTODN35	HTOD045	HTOEA53	HTOEB55	HTOEB76
Gene No.	2379	2380	2381	2382	2383	2384	2385	2386	2387	2388

Last AA of ORF	24	24	46	43	23	40	14	37	15	31	71
First AA of Secreted Portion			8	19		20		32		30	46
Last AA of Sig Pep			L	18		19		31		29	45
First Last AA AA of of Sig Sig Pep Pep	Ţ	1	1	1	1	1		1	1	-	1
AA SEQ ID NO: Y	4996	4997	4998	4999	2000	5001	5002	5003	5004	5005	2006
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	518	518	194	66	159	138	358	922	98	100	26
5' NT of Start Codon	518	518		66		138		922	98	100	26
	1897	1897	2206	2597	1075	2778	1904	1918	1768	2196	1561
5' NT of Clone Seq.	434	434	1	1	1			773			1
Total NT Seq.	1921	1920	2206	2597	1075	2778	1904	1918	1768	2196	1561
NT SEQ D NO:	2399	2400 1920	2401 2206	2402	2403 1075	2404 2778	2405 1904	2406 1918	2407 1768	2408	2409 1561
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203917	203917 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203959 04/26/99
cDNA Clone ID	HTOET03	нтоет03	HTOEV01	HTOFA11	НТОРСЗЗ	HTOGB79	HTOHE22	HTOHG63	нтон193	HTOHM12	HTOHM82
Gene No.	2389	2390	2391	2392	2393	2394	2395	2396	2397	2398	2399

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Last AA of ORF	34	37	14	36	35	23	30	47	134	48
First AA of Secreted Portion	22	19		26	24	20	21	20	44	17
Last AA of Sig Pep	21	18		25	23	19	20	19	43	16
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	1	-	1	-	1	1	1
	5007	2008	2009	5010	5011	5012	5013	5014	5015	5016
S' NT AA of SEQ First ID AA of NO: Signal Y Pep	50	110	275	56	66	160	215	296	107	1034
5' NT of Start Codon	50	110	275	56	66		215	296	107	1034
3' NT of Clone Seq.	1963	1300	1104	1472	1117	1797	1435	1472	1447	3003
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1	-	Ţ		П	Н		-	-	838
Total NT Seq.	1963	1300	1146	1472	1117	1797	1435	1472	1447	3003
NT SEQ D NO:	2410	2411	2412	2413	2414	2415	2416	2417	2418 1447	2419 3003
Vector	Uni-ZAP XR									
ATCC Deposit No.Z and Date	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203959 04/26/99
cDNA Clone ID	HTOHN40	HTOHR59	HTOHS29	HTOID65	HTOE17	HTOIG16	НТОІН39	HTOIH51	HTOJB02	HTOJJ26
Gene No.	2400	2401	2402	2403	2404	2405	2406	2407	2408	2409

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Last AA of ORF	33	53	33	33	53	40	09	31	47	13
First AA of Secreted Portion	19	19	27	21	29	24	36	25	19	
Last AA of Sig Pep	18	18	26	20	28	23	35	24	18	į
AA First Last SEQ AA AA ID of of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	-	1
	5017	5018	5019	5020	5021	5022	5023	5024	5025	5026
5' NT of of of Start AA of Codon Signal Pep	268	130	244	234	145	73	171	509	240	153
5' NT of Start Codon	268		244	234	145		171	509	240	153
3' NT of Clone Seq.	1524	1842	1895	1641	1445	1467	1293	2068	389	1573
S' NT 3' NT of Of Clone Seq. Seq.	-	-	-	189	134	-	Н	175	-	1
Total NT Seq.	1524	1842	1895	1641	1807	1467	1293	2068	389	2027
NT SEQ D NO:	2420	2421	2422	2423	2424	2425	2426	2427	2428	2429 2027
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pBluescript
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	PTA- 181 06/07/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203979 04/29/99
cDNA Clone ID	HTOJP25	HTOJS23	HTOJY56	HTOJZ18	HTPCG10	HTPCO75	HTPCW21	HTPDD68	HTPDV75	HTSER28
Gene No.	2410	2411	2412	2413	2414	2415	2416	2417	2418	2419

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Last AA of ORF	17	20	49	21	35	45	15	37	66	234
First AA of Secreted Portion	16	42	10		34	31		33	18	14
Last AA of Sig Pep	15	41	6		33	30		32	17	13
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	1	1	1	1	1	1	1	1	1
AA SEQ ID Y	5027	5028	5029	5030	5031	5032	5033	5034	5035	5036
5' NT of First AA of Signal	1	134	669	249	39	104	282	130	277	1024
5' NT of of of Start AA of Codon Signal	1147	134	669	249	68	104	285	130		
3' NT of Clone Seq.	1345	1093	879	862	1050	1040	2329	524	711	1992
5° NT 3° NT of of Total Clone Clone NT Seq. Seq.	896		337		1	15	401	99	1	856
Total NT Seq.	1345	1093	1300	862	1050	1040	2364	524	711	1992
NT SEQ ID NO:	2430	2431	2432	2433	2434	2435	2436 2364	2437	2438	2439
Vector	pBluescript	pBluescript	pBluescript	pBluescript	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203959 04/26/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203959 04/26/99	203918 04/08/99	203959 04/26/99
cDNA Clone ID		HTSFV18	HTSG013	HTSG088	HTTAH05	HTTAP37	HTTBJ38	HTTDB11	HTTDG27	HTTDN24
Gene No.	2420	2421	2422	2423	2424	2425	2426	2427	2428	2429

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Last AA of ORF	244	89	57	7	40	78	27	47	32	9/
First AA of Secreted Portion	31	33	31	9	21	21	20		23	44
Last AA of Sig Pep	30	32	30	5	20	20	19		22	43
First AA of Sig Pep			П	-	-			-	-	1
AA SEQ ID NO: Y	5037	5038	5039	5040	5041	5042	5043	5044	5045	5046
		44	188	236	1133	52	30	160	111	137
5' NT of of of Start AA of Codon Signal Pep	22	44			1133	52	30	160	111	137
3' NT of Clone Seq.	1161	1255	2204	1409	2379	1338	1081	1877	1332	773
5' NT 3' NT of of Clone Clone Seq. Seq.	_	-		-	1084	-	Н	-	-	-1
Total NT Seq.	1161	1255	2204	1409	2389	1338	1081	1877	1352	773
NT SEQ ID NO:	2440	2441	2442	2443	2444	2445	2446 1081		2448	2449
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Lambda ZAP II 2447	pSport1	pSport1
ATCC Deposit No.Z and Date	PTA- 1838 05/09/00	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203918 04/08/99	203918 04/08/99	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HTTD033	HTTDT67	HTTE025	HTTEP11	HTTES77	HTTFD29	HTTFG15	HTWAM19	HTWBF58	HTWB030
Gene No.	2430	2431	2432	2433	2434	2435	2436	2437	2438	2439

Last AA of ORF	28	50	54	16	318	24	33	57	32	46
First I AA of Secreted Portion C	28	18	30		23	17	27	14	21	30
	27	17	29		22	16	26	13	20	29
First AA of Sig Pep	1	-	-	1	1	1	-	1	_	
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	5047	5048	5049	5050	5051	5052	5053	5054	5055	5056
F 5' NT AA First of SEQ AA of First ID of Start AA of NO: Sig Codon Signal Y Pep Pep	066	244	892	143	302	244	198	132	133	93
5' NT of Start Codon	066	244	892	143	302	244	198		133	93
3' NT of Clone Seq.	1658	1241	1036	1560	1390	1472	863	1066	1436	613
5' NT 3' NT of of Clone Clone NT Seq. Seq.	777	_	828				П	Н		83
Total NT Seq.	1667	1241	1054	1560	1390	1472	893	1066	1436	684
NT SEQ ID NO:	2450	2451	2452	2453	2454 1390	2455 1472	2456	2457	2458	2459
Vector	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	203959	203918 04/08/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203959	203918 04/08/99	203918 04/08/99
cDNA Clone ID	HTWBZ57	HTWCC10	HTWCE14	HTWCT76	HTWDJ17	HTWDM89	HTWEA05	HTWEG06	нтwеQ36	HTWFA21
Gene No.	2440	2441	2442	2443	2444	2445	2446	2447	2448	2449

Last AA of ORF	116	39	30	32	20	41	26	16	34	15
First Secreted Portion (14	21	21	29	30	20	16		19	
	13	70	70	28	59	19	15		18	
First Last AA AA of of Sig Sig Pep Pep	1	-		-	1	1	1	1	1	1
AA SEQ ID NO: Y	5057	5058	5059	5060	5061	5062	5063	5064	5905	5066
	293	680	193	200	30	49	35	261	159	153
5' NT of of of Start AA of Codon Signal Pep	293	089	193	200	30		35	261	159	
3' NT of Clone Seq.	1851	1693	1298	504	761	1924	1600	759	176	1573
5' NT of Clone Seq.	1	319	1	_	1	_	1		1	1
Total NT Seq.	1851	1693	1298	504	761	1924	1600	759	176	1573
SEQ BD NO:	2460	2461	2462	2463	2464	2465 1924	2466 1600	2467	2468	2469 1573
Vector	pSport1	pSport1	pSport1	Lambda ZAP II 2463	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	PTA- 181 06/07/99	203959 04/26/99	203959	203918	203918 04/08/99	203918 04/08/99	PTA- 1838 05/09/00
cDNA Clone ID	HTWFA88	HTWFM85	HTWF043	HTWLG39	HTXAA20	HTXAD75	HTXAR92	HTXBS38	HTXBU88	HTXCP27
Gene No.	2450	2451	2452	2453	2454	2455	2456	2457	2458	2459

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Last AA of ORF	14	64	42	45	27	31	99	70	34	27	43
First AA of Secreted Portion		23	28	17	26	30	47	11	27	18	33
Last AA of Sig Pep		22	27	16	25	29	46	10	26	17	32
First AA of Sig Pep	-	-	1	1	1	1	1	-	-	1	1
AA SEQ ID NO:	2067	2068	5069	5070	5071	5072	5073	5074	5075	5076	5077
5' NT of First AA of Signal Pep		181	189	253	210	118	149	267	168	323	151
5' NT of Start Codon		181		253		118	149	267	168	323	151
3' NT of Clone Seq.	1440	1544	1335	1628	1957	369	1320	1521	1103	896	1544
5' NT 3' NT of of Clone Clone NT Seq. Seq.	86	1	15	1	1	-		1	1	-	1
Total NT Seq.	1440	1544	1335	1628	1957	636	1320	1521	1103	896	1544
NT SEQ ID NO:		2471	2472	2473	2474	2475	2476	2477	2478	2479	2480
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203959 04/26/99	203918 04/08/99	203918 04/08/99	203959	203959	203959 04/26/99	203959 04/26/99	203959	203918 04/08/99
cDNA Clone ID	HTXCU30	HTXCV44	HTXDJ21	HTXDJ75	HTXDJ85	HTXDK09	HTXD017	HTXDT72	HTXDU08	HTXDZ68	HTXEN33
Gene No.	2460	2461	2462	2463	2464	2465	2466	2467	2468	2469	2470

Last AA of ORF	81	15	32	32	4	31	83	12	6
First I AA of Secreted Portion C	18			31		17	23	7	
	17			30		16	22	9	
first AA of Sig Pep	-	—		1	1	1	-		_
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	5078	5079	5080	5081	5082	5083	5084	5805	5086
	1	132	102	113	192	211	204	1809 5085	275
S' NT of of Of Start AA of Codon Signal Pep	179	132	102	113	192	211	204		275
3' NT of Clone Seq.	<i>LL</i> 9	1678	1393	1425	1238	1117	1640	1195	823
S' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1	-	-		1	1		49	1
Total NT Seq.	219	1678	1679	1425	1238	1117	1640	2060	823
SEQ NO:	2481	2482 1678	2483	2484 1425	2485	2486 1117	2487 1640	2488 2060	2489
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	PTA- 1838 05/09/00	PTA- 1838 05/09/00	203917	203918 04/08/99	203959 04/26/99	PTA- 181 06/07/99	PTA- 791 09/27/99	PTA- 793 09/27/99
cDNA Clone ID	HTXES13	HTXFD86	HTXGK12	HTXGL32	HTXJD08	HTXJD85	HTXJE12	HTXJIS9	HTXJJ92
Gene No.	2471	2472	2473	2474	2475	2476	2477	2478	2479

F A St	m T		129		12	46	52	46	38	55
	73	6	- 12	61		4	2	4	6	
First AA of Secreted Portion	47		20	17		22	27	25	23	21
	46		19	16		21	76	24	22	20
First AA of Sig Pep	_	1	1	1	1	-	-	-	1	-
	5087	2088	5089	5090	5091	5092	5093	1245 5094	5095	5096
		569	635	<i>L</i> 98	350	153	54	1245	133	286
5' NT of of of Start AA of Codon Signal Pep	44	269	635	<i>L</i> 98		153	54	1245	133	286
	938	1896	1494	2801	1073	1290	1629	1610	1945	1455
S' NT 3' NT of Clone Seq.	1	_	511	570	1		1	278		-
Total NT Seq.	938	1896	1494	2836	1073	1290	2496 1629	2497 1610	1945	2499 1455
SEQ NO:	2490	2491	2492 1494	2493 2836	2494	2495 1290	2496	2497	2498	2499
Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203918	203979	203979	203918	203918	PTA- 181 06/07/99	203959	203918	PTA- 792 09/27/99
cDNA Clone ID	HTXJM94	HTXJV54	HTXJW06	HTXKB57	HTXKH22	HTXKH40	HTXKK76	HTXKL53	HTXKS11	HTXKS27
Gene No.	2480	2481	2482	2483	2484	2485	2486	2487	2488	2489

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Last AA of ORF	08	20	13	35	91	14	26	4	92	35
Last First of AA of Sig Secreted Pep Portion	29	23		25	10		17	22	24	16
Last AA of Sig Pep	28	22		24	6		16	21	23	15
First AA of Sig Pep	1	-	1	1	1	1	-	-	-	-
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	2097	2098	5099	5100	5101	5102	5103	5104	5105	5106
	295	359	189	893	674	272	1070	006	93	44
5' NT of of of Start AA of Codon Signal Pep	295	359	189	893		272	1070	006	93	44
3' NT of Clone Seq.	743	715	1038	1568	2058	840	2372	1458	1127	772
5' NT 3' NT of of Clone Clone NT Seq. Seq.	1	1	1	869	1	1	627	559	-	-
Total NT Seq.	743	715	1040	3511	2058	840	2387	2507 2064	1127	772
SEQ NO:	2500	2501	2502	2503 3511	2504 2058	2505	2506 2387	2507	2508	2509
Vector	Uni-ZAP XR	pBluescript								
ATCC Deposit No.Z and Date	203959	203959	203959	203959	203959	203959	203959	203959	PTA- 1838 05/09/00	203918 04/08/99
cDNA Clone ID	HTXLC05	HTXLC45	HTXLT36	HTXLY94	HTXNV66	HTXOL30	HTXOW27	HTXPD86	HTXPT57	HTYSJ88
Gene No.	2490	2491	2492	2493	2494	2495	2496	2497	2498	2499

Last AA of ORF	39	39	38	32	73	13	328	31	16	109	31
First I AA of AS of Portion C	34	37	18	14	13		24			23	24
AA of Sig Pep	33	36	17	13	12		23			22	23
First I AA of Sig Pep	1	1	1	-	1	_	1	1	_	1	-
AA SEQ ID NO:	5107	5108	5109	5110	5111	5112	5113	5114	5115	5116	5117
	202	147	232	54	304	229	09	1041	174	95	453
5' NT of of of Start AA of Codon Signal Pep	202	147	232	54	304	229	09	1041	174	96	453
	1014	1642	1534	857	819	739	1537	2130	1384	1374	743
S' NT 3' NT of of Clone Clone Seq.	-	1		-	1	1		887	1		-
Total NT Seq.	1014	1642	1534	857	819	739	1537	2146	1384	1374	743
SEQ N	2510 1014	2511	2512 1534	2513	2514	2515	2516 1537	2517	2518	2519	2520
Vector	ZAP Express	ZAP Express	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203918	203918 04/08/99	203918 04/08/99	203959	203918 04/08/99	203959 04/26/99	203959	203918 04/08/99	203959 04/26/99
cDNA Clone ID	HUDBE20	HUDBK47	HUFAB57	HUFAL17	HUFA092	HUFA094	HUFAP33	HUFAU71	HUFBK95	HUFBP77	HUFBV62
Gene No.	2500	2501	2502	2503	2504	2505	2506	2507	2508	2509	2510

Last AA of ORF	25	30	20	50	32	9	232	232	36	28	23
First Last AA of AA Secreted of Portion ORF	20	16		44	16		21	21	18	21	19
	19	15		43	15		20	20	17	20	18
AA First Last SEQ AA AA Of Of Of Sig Sig Y Pep Pep	-	-	-	-		1	1	1	-		
AA SEQ ID NO: Y	5118	5119	5120	5121	5122	5123	5124	5125	5126	5127	5128
	334	253	201	200	61	7815	201	201	63	94	220
5' NT of of of Start AA of Codon Signal Pep		253	201	200	61	815	201	201	63	94	
	736	803	1010	1554	1700	2058	1773	1773	275	646	529
S' NT 3' NT of Clone Seq.	1	1	-	1		730	32	32	-	-	
Total NT Seq.	736	803	1010	1554	1700	2058	1781	1781	575	646	529
SEQ NO:	2521	2522	2523	2524	2525	2526	2527	2528	2529	2530	2531
Vector	pSport1	pSport1	pSport1	Lambda ZAP II 2524	Lambda ZAP II 2525 1700	Lambda ZAP II 2526	Lambda ZAP II 2527 1781	Lambda ZAP II	Lambda ZAP II 2529	Lambda ZAP II 2530	Lambda ZAP II 2531
ATCC Deposit No.Z and Date	203959	203918 04/08/99	203959	203918 04/08/99	203979	203959	203959	203959	203959 04/26/99	203959 04/26/99	203959 04/26/99
cDNA Clone ID	HUFBY96	HUFCN72	HUFEF79	HUKAD46	HUKAI28	HUKA050	HUKCS86	HUKCS86	HUKEA22	HUKEL79	HUKEX37
Gene No.	2511	2512	2513	2514	2515	2516	2517	2518	2519	2520	2521

Last AA of ORF	6	6	477	13	10	486	300	000	510	16	37
First I. AA of A Secreted Portion C.			46			2	71		39		20
		!	45			1	15	CI	38		19
First AA of Sig Pep	-	-	_	-	-		- 1	-	—	1	-
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	5129	5204	5130	5131	5132	5133	70,7	5134	5135	5136	5137
	260		49	237	830	3	t	297	101	205	168
5' NT of of of Start AA of Codon Signal Pep	260	261	49	237	830					205	168
	696	973	1531	957	1121	1664	ļ	1664	1986	1340	718
S' NT 3' NT of of Clone Clone Seq. Seq.	1	1	20	1	1	24		24	П	-	-
Total NT Seq.	963	1177	1574	2735	1121	1971		1971	1986	1340	719
SEQ X	2532	7097		2534	2535	2536			2538	2539	2540
Vector	Lambda ZAP II 2532	Lambda ZAP II 2607	Lambda ZAP II 2533	Lambda ZAP II 2534 2735	pBluescript SK- 2535	Lambda ZAP II 2536 1971		Lambda ZAP II 2537	Lambda ZAP II 2538 1986	Lambda ZAP II 2539	pSport1
ATCC Deposit No.Z and Date	203979	+			203918		791 09/27/99	PTA- 791 09/27/99	PTA- 181	203918	203959
cDNA Clone ID	HUKFC71	HUKFC71	HUKFV37	HUKFY09	HUNAL39	HUSA004		HUSA004	HUSCA09	HUSCJ01	HUSGB23
Gene No.	2522	2522	2523	2524	2525	2526		2527	2528	2529	2530

Last AA of ORF	32	34	22	72	487	33	41	56	26	52
First AA of Secreted Portion (24			20	22	21	21	23	24	30
Last AA of Sig Pep	23			19	21	20	20	22	23	29
	1	-	-	1	1	1	1	1	1	1
AA First SEQ AA ID of NO: Sig Y Pep	5138	5139	5140	5141	5142	5143	5144	5145	5146	5147
5' NT of SEQ of First ID Start AA of NO: Codon Signal Y	191	73	643	227	812	204	177	301	281	42
5' NT of Start Codon	191	73	643	227	812	204			281	42
	889	1940	1526	2576	2758	1415	925	669	1236	413
5' NT 3' NT of of Of Clone Clone NT Seq. Seq.	1	1	299	1	768	1	-	1	1	1
Total NT Seq.	889	1940	1526	2576	6705	2546 1415	925	669	1236	413
X SEQ X	2541	2542	2543	2544	2545	2546	2547	2548	2549	2550
Vector	pSport1	pSport1	pSport1	Lambda ZAP II 2544 2576	Lambda ZAP II 2545	pSport1	pSport1	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	203918 04/08/99	203918 04/08/99	203918 04/08/99	PTA- 181 06/07/99	203979 04/29/99	PTA- 181 06/07/99	PTA- 793 09/27/99	203918 04/08/99	203959 04/26/99	203918 04/08/99
cDNA Clone ID	HUSG109	HUSGQ57	HUSGY15	HUSHD41	HUSHK65	HUSIK45	HUSIO57	HUSIP17	HUSIR70	HUSXP50
Gene No.	2531	2532	2533	2534	2535	2536	2537	2538	2539	2540

Last AA of ORF	19	69	25	38	22	92	87	25	31	15
First AA of Secreted Portion		29	18	29	13	40	27		16	
		28	17	28	12	39	26		15	
First AA of Sig Pep	-	-		-	1	1	1	-	-	-
AA] SEQ ID NO: Y	5148	5149	5150	5151	5152	5153	5154	5155	5156	5157
	148	421	265	27	366	479	443	488	258	318
5' NT of of of Start AA of Codon Signal Pep	148	421	265	27		479	443	488	258	318
3' NT of Clone Seq.	639	744	829	2785	2163	1278	1649	1727	2314	1161
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1	363	T	-	151	1	34	464	-	
Total NT Seq.	639	744	8/9	2785	2555 2163	1278	1649	1727	2314	1161
SEQ NO:	2551	2552	2553	2554 2785	2555	2556	2557	2558	2559	2560
Vector	pSport1	pSport1	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 3.0 2556	pCMVSport 3.0 2557 1649	pCMVSport 3.0 2558	pCMVSport 3.0 2559	pCMVSport 3.0 2560 1161
ATCC Deposit No.Z and Date	203918 04/08/99	203959	203918 04/08/99	PTA- 1838	203959	203959	203959	3918	203959	203959 04/26/99
cDNA Clone ID	HUSXY93	HUSYG26	ниусое8	HUVDG58	HUVEG53	HWAAH11	HWAAQ28	HWAAY60	HWABR43	НWАСН06
Gene No.	2541	2542	2543	2544	2545	2546	2547	2548	2549	2550

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Last AA of ORF	62	22	405	17	51	9	54	30	4
First AA of Secreted Portion	23		24	16	23	33	37	29	32
Last AA of Sig Pep	22		23	15	22	32	36	28	31
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	1	-	1	1	1	-	1	1	1
	5158	5159	5160	5161	5162	5163	5164	5165	5166
	110	314	212	260	783	615	140	261	134
5' NT of of of Start AA of Codon Signal Pep		314	212	260	783	615	140	261	134
3' NT of Clone Seq.	1462	2393	2193	372	2731	2783	625	831	1468
of of Clone Seq.	1	1	-	П	623	1	1	_	1
Total NT Seq.	1462	2393	2193	372	2731	2783	625	831	1468
SEQ NO:		2562		2564	2565		2567	2568	2569
Vector	pCMVSport 3.0 2561	pCMVSport 3.0 2562 2393	pCMVSport 3.0 2563	pCMVSport 3.0 2564	pCMVSport 3.0 2565	pCMVSport 3.0 2566	pCMVSport 3.0 2567	pCMVSport 3.0 2568	pCMVSport 3.0 2569
ATCC Deposit No.Z and Date	203959 04/26/99	PTA- 1838 05/09/00	PTA- 792 09/27/99	PTA- 181 06/07/99	66/63	PTA- 181 06/07/99	203959 04/26/99	PTA- 181 06/07/99	203979 04/29/99
cDNA Clone ID	HWACZ33	HWADV90	HWAEB52	HWBAK71	HWBBU75	HWBCN81	HWBCP16	HWBCX93	HWBEF34
Gene No.	2551	2552	2553	2554	2555	2556	2557	2558	2559

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Last AA of ORF	38	17	34	61	61	4	29		39	71
First AA of Secreted Portion	30	i	18	24	48	29	21		29	24
	29		17	23	47	28	70		78	23
First Last AA AA of of Sig Sig Pep	-	1	1	-	1	1	1		-	-
	5167	5168	5169	5170	5171	5172	5173	5174	5175	5176
5' NT AA 5' NT of SEQ of First ID Start AA of NO: Codon Signal Y		315	099	268	213	166	148	2741	42	1091
5' NT of Start Codon	40	315	099	268	213	166		2741	42	1091
	1411	875	1026	596	969	871	843	2932	536	1835
5' NT 3' NT of of Clone Seq. Seq.	1	1	213	1	1	1	1	2488	-1	1007
Total NT Seq.	1411	875	1026	596	695	871	843	2973	536	2579 1898
SEQ NO:	2570	2571	2572	2573	2574	2575	2576	2577	2578	2579
Vector	pBluescript SK- 2570 1411	pBluescript SK- 2571	pCMVSport 3.0 2572	pCMVSport 3.0 2573	pCMVSport 3.0 2574	pCMVSport 3.0 2575	pCMVSport 3.0 2576	pSport1	pSport1	pSport1
ATCC Deposit No.Z and Date	PTA- 181 06/07/99	203918 04/08/99	203959	203959	203959	203959	PTA- 181 06/07/99	203959 04/26/99	203959	203959 04/26/99
cDNA Clone ID	HWFBB23	HWFBI40	HWHGV77	HWHGW09	НWННА21	HWHPU44	HWHRC51	HWLAT50	HWLBO67	HWLGP26
Gene No.	2560	2561	2562	2563	2564	2565	2566	2567	2568	2569

Last AA of ORF	10	39	31	20	74	13	10	44	33	38
First I AA of Secreted Portion C		32	24		18			56	22	
		31	23		17			25	21	
First AA of Sig Pep	1	1	1	1	1	1	1	-		-
AA First Last SEQ AA AA ID of of NO: Sig Sig Y Pep Pep	5177	5178	5179	5180	5181	5182	5183	5184	5185	5186
S' NT AA of SEQ First ID AA of NO: Signal Y	183	531	594	2526 5180	377	1523 5182	33	228	197	2270 5186
5' NT of of Of Start AA of Codon Signal	183		594	2526	377			228	197	2270
3' NT of Clone Seq.	1701	787	1030	2770	598	2306	91	669	338	2789
5' NT 3' NT of of Clone Clone NT Seq. Seq.	-	-		2583 2770 2287	245	1432	1		1	2179
Total NT Seq.	1701	787	1030	2770	598	2585 2306	91	669	338	2589 2789
NT SEQ ID T NO:	2580	2581	2582	2583	2584	2585	2586	2587	2588	2589
Vector	pSport1	pSport1	pSport1	pSport1	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959	203959	203959	203959	203918 04/08/99	203959	203959	PTA- 181 06/07/99	203918	203959 04/26/99
cDNA Clone ID	НWLНО31	HWLIL31	HWLJN08	HWLRE03	HWTAM38	HWTAW58	HWTBB42	HWTBC75	HWTB125	HWTBL86
Gene No.	2570	2571	2572	2573	2574	2575	2576	2577	2578	2579

+ , <u>F</u>								
Last AA of ORF	271	38	18	31	37	89	20	22
First AA of Secreted Portion	12	19		18	27	34	19	12
	11	18		11	26	33	18	11
First AA of Sig Pep	1	1		-	-	-	1	-
AA SEQ D D NO:	5187	5188	5189	5190	5191	5192	5193	5194
5' NT of First AA of Signal Pep	261	197	341	461	166	261	116	151
		197	341	461		261	116	151
3' NT of Clone Seq.	1145	592	2230	1223	1168	1193	584	968
5' NT 3' NT of of Ot Clone Clone NT Seq. Seq.	1	1	1	1	-	-	1	-
	1145	592	2230	1223	1168	1193	584	968
NT SEQ ID NO:		2591	2592	2593	2594 1168	2595	2596	2597
Vector	Uni-ZAP XR 2590	pCMVSport 3.0 2591	pCMVSport 3.0 2592	pCMVSport 3.0 2593	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC Deposit No.Z and Date	203959 04/26/99	203918 04/08/99	PTA- 1838 05/09/00		PTA- 181 06/07/99	PTA- 1838 05/09/00		203918 04/08/99
cDNA Clone ID	HWTBX66	HYAAC74	HYAAD61	HYACC21	HYBAP75	HYBAQ24	HYBAW56	HYBBD81
Gene No.	2580	2581	2582	2583	2584	2585	2586	2587

TABLE 1B	E 1B			:			i i i i i i i i i i i i i i i i i i i		
7	1	2,445	or oas	ORE	AA SEO		Tissue Distribution Library code: count	Cytologic	OMIM
Sene No:	Clone ID NO: Z	Conug D:	NO: X	(From-To)	NO.	Predicted Epitopes	(see Table 4 for Library Codes)	Band	Disease Reference(s):
-	HJCRDJU	907073	=	27 - 80	2608	Gln-1 to Asp-6.	T0110: 1 and L0776: 1.		
2	H2CBH91	826669	12	309 - 431	2609	Met-1 to Lys-12.	AR089: 3, AR316: 3,	8p22-q21.2	148370, 238600, 238600, 238600,
							L0764: 3, L0746: 2, L0749:		238600, 600143,
							2, L0777: 2, S0114: 1, H0661:		601385, 602629
							1, H0497: 1, H0036: 1, T0110:		
							1, H0412: 1, L0520: 1, L0651: 1 1 0796: 1 1 0373: 1, L0766:		
							1, L0806: 1, L0776: 1, L0789:		
							1, L0666: 1, L0742: 1, L0745:		
							1, L0731: 1, L0485: 1 and		
						,	L0608: 1.		,,0,0,
7	H2I BAS4	684290	13	386 - 403	2610		L0766: 5, L0439: 5, L0803:	6q21	120110, 121014,
n	TCAGTZII	077	}				4, L0666: 4, H0556: 3, S0360:		601666, 602772
	_						3, H0591: 3, L0809: 3, L0754:		
							(3, L0750: 3, L0777: 3, H0392:		
							2, H0553: 2, L0771: 2, L0662:		
							2, L0794: 2, L0806: 2, L0748:		
							2, L0749: 2, L0779: 2, L0759:		
							[2, H0707: 2, S0026: 2, H0171:		
							1, H0450: 1, H0125: 1, S0376:		
							1, H063/: 1, H0380: 1, S0043:		
							11, S0222: 1, H0451: 1, H0497:		
							1, H0599: 1, H0581: 1, T0115:		
						-	1, L0471: 1, H0051: 1, S0022:		
<u> </u>							1, L0142: 1, H0674: 1, H0038:		
							1, H0634: 1, H0616: 1, S0386:		
·		****					1, H0625: 1, H0646: 1, L0804:		
							1, L0774: 1, L0805: 1, L0655:		
							1, L0559: 1, L0526: 1, L0790:		
							1, L0663: 1, H0519: 1, H0593:		
<u>.</u>			-				1, H0684: 1, H0659: 1, H0670:	••	
							1, L0/55: 1, L0/51: 1, H0445:		

	182600, 232700, 602086	182600, 232700, 602086
	14q21.1-q21.3	14q21.1-q21.3
1, L0605: 1, L0599: 1, L0608: 1, L0601: 1, H0543: 1 and S0424: 1.	AR089: 5, AR316: 5, AR060: 4 L0773: 7, S0410: 6, L0803: 6, L0752: 5, S0358: 4, L0766: 4, L0439: 4, L0646: 3, L0659: 3, S0442: 2, S0408: 2, S0003: 2, H0038: 2, S0408: 2, L0564: 2, L0655: 2, L0666: 2, L0664: 2, L0655: 2, H0521: 2, L0748: 2, L0740: 2, L0777: 2, L0731: 2, L0758: 2, L0362: 2, H0170: 1, S0116: 1, H0402: 1, S0444: 1, S0360: 1, S0476: 1, L0717: 1, H055: 1, H0590: 1, S0346: 1, H0050: 1, S0051: 1, H0529: 1, H0412: 1, L0731: 1, H0529: 1, L0483: 1, H0531: 1, H0529: 1, L0761: 1, L0772: 1, L0794: 1, L0807: 1, S0374: 1, H0435: 1, H0539: 1, H0518: 1, S0152: 1, H0539: 1, H0518: 1, S0152: 1, H0423: 1 and H0506: 1.	AR089: 5, AR316: 5, AR060: 4 L0773: 7, S0410: 6, L0803: 6, L0752: 5, S0358: 4, L0766: 4, L0439: 4, L0646: 3, L0659: 3, S0442: 2, S0408: 2, S0003: 2, H0038: 2, S0440: 2, L0774: 2, L0655: 2, L0666: 2, L0664: 2, L0740: 2, L0777: 2, L0731: 2, L0758: 2, L0362: 2, H0170: 1, S0116: 1, H0402: 1, S0444: 1, S0360: 1, S0476: 1, L0717:
	Lys-28 to Thr-33.	Lys-28 to Thr-33.
	2611	2612
	197 - 307	236 - 346
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		830636
	H2LBB09	H2LBB09
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1, H0574: 1, H0632: 1, T0109: 1, H0156: 1, H0590: 1, S0346: 1, H0052: 1, T0115: 1, H0597: 1, H0050: 1, S0051: 1, H0622: 1, L0483: 1, H0553: 1, H0622: 1, H0412: 1, L0351: 1, H0529: 1, L0761: 1, L0772: 1, L0794: 1, L0807: 1, S0374: 1, H0435: 1, H0539: 1, H0518: 1, S0152: 1, S0392: 1, S0436: 1, H0653: 1, H0423: 1 and H0506: 1.	3, L0794: 2, H0458: F: 1, H0617: 1, L0772: F: 1, L0659: 1, L0663: 7758: 1.	AR270: 8, AR061: 7, AR060: 5, AR238: 5, AR269: 5, AR268: 5, AR296: 4, AR290: 4, AR233: 4, AR290: 4, AR291: 4, AR293: 4, AR291: 4, AR183: 4, AR291: 4, AR298: 4, AR291: 4, AR298: 4, AR291: 4, AR298: 4, AR299: 3, AR291: 1, AR299: 1, AR291: 1, AR295: 1, AR291: 1, AR293: 2, AR291: 2, S0250: 2, L0764: 2, L0794: 2, L0381: 2, H0556:
	2613	2614
	172 - 270	717 - 1055
	16	17
	610045	826161
	H2MAC63	H2MBA76
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	126060, 143200, 143200, 181510, 214300, 253200, 600354, 600354, 600387
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	2616	2617
	95 - 217	196 - 936
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2, H0690: 2, H0555: 2, L0743: 2, L0755: 2, L0757: 2, L0759: 2, S0436: 2, L0588: 2, H0352: 2, H0624: 1, H0685: 1, H0295: 1, S0134: 1, H0683: 1, H0686: 1, H0341: 1, S0134: 1, H0341: 1, S0134: 1, H0341: 1, S0134: 1, H0696: 1, S0448: 1, H0339: 1, H0609: 1, H0019: 1, H0037: 1, H0023: 1, H0069: 1, L0770: 1, L07	L0740: 8, L0748: 7, L0749: 6, L0805: 4, L0438: 4, S0426: 3, L0766: 3, L0752: 3, H0657: 2, S0474: 2, L0804: 2, L0659: 2, L0809: 2, L0754: 2, L0439: 2, L0754: 2, L0592: 2, L0592: 2, L0508: 2, S0202: 2, L0592: 2, L0608: 2, S0202: 2, L0592:
4444	
•	2621 Gln-21 to Gly-26, Cys-30 to Ser-38.
	24 86 - 229
	845956
	14 H7TDB54

1, 70002: 1, S0040: 1, S0444: 1, S0360: 1, T0008: 1, H0329: 1, H0592: 1, H0486: 1, L0021: 1, H0123: 1, H0373: 1, H0266: 1, H0188: 1, S0003: 1, S0214: 1, H0553: 1, H0090: 1, H0591: 1, H0551: 1, S0438: 1, L0772: 1, L0372: 1, L0662: 1, L0794: 1, L5564: 1, L0803: 1, L0774: 1, L0555: 1, L0527: 1, L0783: 1, L0790: 1, L0666: 1, H0658: 1, H0522: 1, H0696: 1, S0406: 1, S0037: 1, L0751: 1, L0745: 1, L0753: 1, H0445: 1, S0026: 1 and H0352: 1.	2	53: 21, AR213: 20, 40: 19, AR096: 18, 52: 16, AR183: 15, 43: 15, AR312: 14, 41: 11, AR309: 11, AR309: 10, 82: 10, AR240: 9, 41: 11, AR290: 9, 55: 9, AR24: 8, 55: 9, AR24: 8, 57: 8, AR192: 5, 5, AR192: 5, AR198: 4, 60: 4, AR198: 4, 60: 4, AR299: 4, 4, 4, 4, AR299: 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4,
1, 70 1, 80 1, 10 1, 10	S01	AR053: AR240: AR052: AR243: AR265: AR265: AR268: AR268: AR267: AR182: AR267: AR182: AR267: AR175: AR175: AR175: AR175: AR175: AR175: AR209:
	2622	2623
	275 - 352	329 - 643
	25	56
	597427	897933
	H7TMB95	HAAAT06
	15	16

AR231: 3, AR293: 3, AR205: 3, AR299: 3, AR177: 3, AR295: 3, AR249: 3, AR292: 3, AR249: 3, AR292: 3, AR296: 3, AR292: 2, AR089: 2, AR291: 2, AR298: 2, AR298: 2, AR298: 2, AR298: 1, AR298: 2, AR299: 1, AR277: 2, AR299: 1, AR314: 1, AR239: 1, AR219: 1 L0752: 10, L0471: 9, L0751: 9, H0422: 9, L0741: 5, L0641: 5, L0641: 5, L0641: 5, L0641: 5, L0791: 2, L0751: 4, S0114: 3, S0360: 3, H0249: 5, H0341: 2, H0031: 2, L0778: 2, L0778: 2, L0769: 2, L0779:	AR205: 3, AR295: 3, AR179: 3, AR295: 3, AR179: 3, AR295: 3, AR179: 3, AR295: 3, AR296: 3, AR295: 2, AR298: 2, AR296: 2, AR296: 2, LOT99: 2, LOT99: 5, LOT99: 2, LOT99:	3,	£ 6.	3,	3,		2,	îc	,	2,	2,	2.		0 1 0731	3, LU/31.	: 6, HUS26:	: 5, L0766:	: 5, H0543:	:: 4, L0662:	3, S0360:	:: 3, H0135:	: 3, L0759:); 3, H0265:	: 2, H0013:	:: 2, H0050:	: 2, L0639:	: 2, L0774:	: 2, L0384:	: 2, H0144:); 2, H0539:	i: 2, L0750:	: 2, S0134:	5: 1, S0212:	: 1, S0046:): 1, H0393:	: 1, H0486:	3: 1. H0545:
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1, H0354: 1, H0271: 1, H0292: 1, H0284: 1, H0286: 1, H0028: 1, H0615: 1, H0622: 1, T0023: 1, H0553: 1, H0616: 1, H0640: 1, S0208: 1, L0796: 1, L0640: 1, L0644: 1, L0796: 1, L0667: 1, L0654: 1, L0659: 1, L0606: 1, L0654: 1, L0659: 1, L0606: 1, L0666: 1, L0659: 1, L0793: 1, L0666: 1, L0438: 1, H0620: 1, H0619: 1, H0689: 1, H0684: 1, H0631: 1, L0741: 1, L0756: 1, S0260: 1, L0599: 1, L0756: 1, S0260: 1, L0599: 1, L0595: 1, S0242: 1 and H0542: 1.	AR060: 6, AR316: 4, AR089: 3 L0769: 5, L0741: 3, L0756: 3, L0806: 2, L0438: 2, S6022: 1, L0520: 1, L0770: 1, L0792: 1, H0144: 1 and S0260: 1.	H0494: 15, L0747: 11, H0586: 10, H0587: 7, L0809: 5, H0600: 4, L0518: 4, L0653: 3, L0743: 3, H0592: 2, H0264: 2, L0648: 2, L0517: 2, S0328: 2, L0744: 2, H0295: 1, S0356: 1, S0132: 1, H0549: 1, S0280: 1, L0021: 1, L0041: 1, H0050: 1, L0455: 1, H0087: 1, H0379: 1, H0477: 1, S0448: 1, L0764: 1, L0789: 1, L0790: 1, S0350: 1, H0696: 1, S0432: 1, L0731: 1, S0384: 1 and S0456: 1.	AR296: 53, AR269: 43, AR291: 30, AR268: 30, AR298: 30, AR285: 30, AR253: 28, AR192: 28,
I, H0354: 1, H0271: 1, H0284: 1, H0286: 1, H0286: 1, H0622: 1, H0653: 1, H0616: 1, L0351: 1, S0208: 1, L0796: 1, L0654: 1, L0654: 1, L0654: 1, L0654: 1, L0656: 1, L0656: 1, L0656: 1, L0656: 1, L0666: 1, H0666: 1, H0661: 1, H0661: 1, H0661: 1, L0699: 1, H0661: 1, L0699: 1, S0260: 1, L0599: 1, S0260: 1, L0599: 1, S0242: 1 and H0541: 1, S0242: 1	AR060: 6, AR089: 3 L0769: 5,1 3, L0806: 2, 1, L0520: 1,	H0494: 15, L0747: 1 H0586: 10, H0587: 7, 5, H0600: 4, L0518: 4, 3, L0743: 3, H0592: 2, 2, L0648: 2, L0517: 2, 2, L0744: 2, H0295: 1, 1, L0021: 1, H0549: 1, 1, L0455: 1, H0087: 1, 1, L0406: 1, S0448: 1, 1, L0789: 1, L0790: 1, 1, H0696: 1, S0432: 1, 1, S0384: 1 and S0456: 1,	AR296: 53, AR291: 30, AR298: 30, AR253: 28,
	Gly-8 to Ala-13.	Ser-16 to Glu-31.	
			9
	2624	2625	2626
	683 - 742	1060 - 1311	654 - 968
	27	58	29
	634372	797625	603946
·	HACAD42	HACBJII	HACBS86
	17	18	19

																																					-		
1 70 AD270. 27	26, AR286:	24, AR266:	22, AR309:	22, AR295:	20, AR175:	20, AR182:				18, AR089:	17, AR183:	17, AR053:	16, AR316:	16, AR186:	AR313: 15, AR247: 15,	AR060: 15, AR271: 15,	AR251: 15, AR104: 14,	AR275: 14, AR267: 14,				13, AR039:		11, AR204:		11, AR299:	10, AR282:		7, AR061:	7, AR280:	5, AR231:	5, AR219:	4, AR315:			3, AR281:	3, AR233:	3, AR232:	AR314: 2
44	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AF	AF	AF	AF	AF	AF	A	A	AI	A	I. A.	A	[A]	A	A	A
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1, L0809: 1, H0435: 1, H0672: 1, H0555: 1, L0749: 1, L0750: 1, H0445: 1, S0434: 1 and H0506: 1.	AR060: 10, AR316: 9, AR089: 8 H0617: 6, L0758: 6, L0770: 4, L0769: 4, L0662: 3, L0665: 3, L0439: 3, L0748: 2, L0758: 1, H0424: 2, L0748: 2, L0751: 2, L0777: 2, L0757: 2, S0356: 1, H0150: 1, H0051: 1, T0010: 1, H0181: 1, H0028: 1, T0006: 1, L0638: 1, L0764: 1, L0771: 1, L0645: 1, L0764: 1, L0771: 1, L0766: 1, L0804: 1, L0775: 1, L0578: 1, L0806: 1, L0775: 1, L0657: 1, L0809: 1, L0689: 1, L0658: 1, L0769: 1, L0689: 1, L0658: 1, L0769: 1, L0676: 1, L0658: 1, L0769: 1, L06776: 1, L0658: 1, L0769: 1, L06776: 1, L0658: 1, L0769: 1, L0776: 1, L0769: 1, L0769: 1, L0776: 1, L0749: 1, L0750: 1, L0752:	AR060: 3, AR316: 2, AR089: 1 S0280: 1 and L0766: 1.	AR089: 2, AR316: 1 L0744: 8, H0427: 4, H0163: 2, S0116: 1, H0021: 1, H0188: 1, S0366: 1, L0646: 1, L0663: 1, S0378: 1 and L0743: 1.	H0427: 1	AR060: 180, AR316: 122, AR089: 57 H0556: 2, S0218: 1, H0427: 1, L0752: 1 and H0136: 1.	H0427: 1, H0553: 1 and
			Pro-45 to Gln-51.			
	2628	2629	2630	2631	2633	2634
	105 - 215	205 - 294	202 - 357	161 - 244	240 - 251	254 - 352
	3.1	32	33	34	36	37
	832528	561629	745367	604500	683227	601695
	HACBZ73	HACCK29	HADAB60	HADAM31	HADCZ65	HADDC04
	21	22	23	24	72	27

H0674: 1.	H0427: 2, H0556: 1, S0001: 1, H0550: 1 and H0539: 1.	AR192: 92, AR207: 79, AR211: 76, AR245: 65, AR193: 64, AR313: 59, AR196: 54, AR313: 59, AR246: 56, AR196: 54, AR191: 50, AR240: 50, AR240: 50, AR240: 50, AR240: 50, AR210: 48, AR211: 47, AR210: 48, AR211: 47, AR218: 46, AR219: 46, AR219: 47, AR299: 41, AR299: 41, AR299: 41, AR299: 42, AR189: 40, AR299: 41, AR299: 41, AR299: 43, AR264: 39, AR316: 39, AR316: 37, AR316: 37, AR316: 37, AR316: 37, AR291: 37, AR291: 31, AR291: 31, AR291: 31, AR291: 31, AR199: 30, AR234: 30, AR292: 33, AR282: 29, AR298: 29, AR298: 29, AR298: 29, AR298: 20, AR298: 25, AR285: 26, AR181: 27, AR285: 26, AR285: 26, AR181: 27, AR285: 26,
		Pro-109 to Ásp-121.
	2635	
	238 - 357	
	38	
	847115	
	HADDP23	
	28	

	25, AR309:		AR233: 24, AR232: 23,	AR268: 23, AR179: 23,			AR250: 22, AR257: 22,				19, AR294:	19, AR296:	 AR283: 17, AR254: 17,	AR291: 16, AR267: 15,	AR256: 15, AR228: 14,		_	-		9, AR168:						AR217: 5	L0747: 19, L0659: 9, L0764:	8, L0665: 8, H0539: 8, L0758:	8, H0617: 7, L0769: 7, L0731:	6, S0360: 5, S0358: 4, L0586:	4, H0427: 4, H0545: 4, S0344:	4, L0763: 4, L0776: 4, H0661:	3, S0420: 3, S0278: 3, H0549:	3, H0550: 3, S0142: 3, L0770:	3, L0648: 3, L0657: 3, L0783:	3, L0749: 3, L0591: 3, H0171:	2, H0294: 2, T0049: 2, H0657:	, H0484: 2, H0255: 2, S0442:
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2, H0586: 5: 2, H0575: 5: 2, H0597:): 2, S0214: : 2, H0616:	t: 2, S0426: : 2, L0768:	: 2, L0775:	: 2, H0660:	: 2, L0752:	L0759: 2, L0590: 1.0504: 3, 50106:	6: 2, H0556:	:: 1, H0295:	: 1, S0212:	9: 1, H0664:	5: 1, S0376:	T0008: 1, H0580:	7: 1, H0441:	H0609: 1, H03/0. T0039: 1, T0060:	S0280: 1, H0156:	, H0274: 1, S0010:	1, H0581: 1, H0052:	1, H0085: 1, H0232:	1, H0544: 1, H0546: 1, H0011: 1, S0051:	3: 1, H0688:	3: 1, H0033:	31: 1, H0644:	16: 1, L0055:	16: 1, H0090:	38: 1, H0087:	9: 1, L0351:	0: 1, H0509: 4: 1 T 0640:	72: 1, L0373:	3: 1, L0662:
2, S0354: 2, S0222: 2, H0586: 2, H0587: 2, H0486: 2, H0575: 2, H0486: 2, H0575: 2, H0597: 2, H0587: 2, H0587: 2, H0587: 2, H0587: 2, H0587: 2, H0587: 2, H0588: 2, H0587: 2, H05	2, H0024: 2, H0510: 2, S0214: 2, H0553: 2, H0181: 2, H0616:	2, H0059: 2, H0494: 2, S0426: 2, 1,0761: 2, L0771: 2, L0768:	2, L0766; 2, L0774; 2, L0775; 3, 1,0653; 3, 1,0653; 4, 1,0653; 5, 1,0653; 5, 1,0663;	2, L0653; 2, L0511; 2, L0660; 2, L0664; 2, S0126; 2, H0660;	2, S0378: 2, L0779: 2, L0752:	2, L0755: 2, L0759	2, L0393; 2, L0394; 2, 30139; 2, H0542; 2, H0506; 2, H0556;	1, T0002: 1, S0402: 1, H0295:	1, S0114: 1, S0298: 1, S0212:	1, H0483: 1, H0669: 1, H0664:	1, H0638: 1, S0356: 1, S0376:	1, S0444: 1, T0008	<u>.</u> .	1, H0431: 1, H0609: 1 1 H0559: 1 T0039: 1	, _ ;	1, H0002: 1, H027	1, H0318: 1, H058		11, H0231: 1, H0544: 1, H0546:	1, H0046; 1, H0011; 1, 5003; 1, H0355; 1, S0003; 1, H0688;	1, H0428: 1, L0483: 1, H0033:	1, H0424: 1, H0031: 1, H0644	1, L0142: 1, H0606: 1, L0055:	1, H0673: 1, H0316: 1, H0090	1, H0591: 1, H0038: 1, H0087	1, H0100: 1, S0039: 1, L0351:	11, S0352: 1, S0440: 1, H0509:	1, H0046: 1, S0144: 1, E0040: 1, L0040: 1, L0040: 1, L0373:	1, L0372: 1, L0773:
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1, L0375: 1, L0651: 1, L0806: 1, L0652: 1, L0654: 1, L0655: 1, L0542: 1, L0647: 1, L0666: 1, S0428: 1, H0547: 1, H0547: 1, H0547: 1, H0547: 1, H0543: 1, S0328: 1, H0548: 1, L0748: 1, L0756: 1, L0748: 1, L0756: 1, L0757: 1, L0748: 1, L0756: 1, L0588: 1, L0581: 1, S0031: 1, S0031: 1, S0031: 1, L0588: 1, H0543: 1, H0553: 1, H0558: 1, H0558: 1, H0553: 1, H05	AR089: 3, AR316: 2, AR060: 1 L0748: 2, L0749: 2, S0222: 1, H0427: 1, H0596: 1, H0644: 1, L0367: 1 and H0352: 1.	H0427: 2 L0748: 8, L0439: 8, L0769: 5, L0666: 5, L0637: 4, L0375: 4, L0779: 4, S0192: 4, H0486: 3, H0427: 3, T0041: 3, L0763: 3, L0770: 3, L0659: 3, L0777: 3, L0758: 3, L0777: 3, L0758: 3, L0777: 3, L0758: 3, L0794: 2, L0776: 2, L0783: 2, L0769: 2, L0749: 2, L0749: 2, L0759: 2, L0759: 2, L0759: 2, L0759: 2, L0759: 2, L0731: 2, L0759: 2, L0759: 2, L0759: 1, S0276: 1, H0402: 1, S0418: 1, H0662: 1, H0402: 1, S0418: 1, S0408: 1, H0402: 1, S0418: 1, S0408: 1, H0402: 1, L0777: 1, S0408: 1, H0402: 1, H0402: 1, L0777: 1, S0408: 1, H0402:
		Met-1 to Trp-8.
	2637	2639
	230 - 337	154 - 249 201 - 248
	40	41 42
	664477	799507
	HADDR24	HADEY08
	30	32

								157640, 174900,
								10q24
1, H0545: 1, H0081: 1, H0011: 1, H0012: 1, H0024: 1, S6028: 1, H0181: 1, H0135: 1, H0040: 1, H0264: 1, H0494: 1, H0494: 1, S0150: 1, H0649: 1, L0667: 1, L0646: 1, L0649: 1, L0649: 1, L0657: 1, L0657: 1, L0657: 1, L0659: 1, L0809: 1, H0144: 1, S0374: 1, H0547: 1, H0588: 1, H0539: 1, S0152: 1, L0743: 1, L0744: 1, H0668: 1 and H0423: 1.	1 1	H0427: 2	H0427: 3, H0497: 1, H0485: 1, H0036: 1, S0010: 1, H0553: 1, L0803: 1, H0701: 1, H0689: 1, H078: 1	H0427: 3, L0759: 3, H0409: 1, H0333: 1, H0013: 1, H0156: 1, L0105: 1, H0083: 1, H0032: 1, L0770: 1, L0796: 1, L0794:	1, L0766: 1, L0774: 1, L0776: 1, L0754: 1, L0777: 1, L0752: 1, L0757: 1.	L0759: 3, S0222: 2, H0427: 2, L0755: 2, L0757: 2, S0134: 1, S0360: 1, H0409: 1, H0333: 1, H0013: 1, H0156: 1, H0575: 1, H0390: 1, L0105: 1, H0263: 1, L0770: 1, L0776: 1, L076: 1, L076: 1, L076: 1, L0755: 1, L0754: 1, L0755: 1, L0755: 1, L0485: 1 and S0196: 1.	H0427: 1	L0163: 4, T0006: 2, H0716:
	Gly-22 to Val-29.	Gly-22 to Val-29.		Ala-17 to Trp-23, Ser-46 to Thr-51.				
	2640	2641	2642	2643		2644	2645	2646
	169 - 267	169 - 267	94 - 234	377 - 577		212 - 328	24 - 197	147 - 314
	43	44	45	46		47	48	49
	704584	861628	668229	829979		843934	637485	793100
	HADEY22	HADEY22	HADFB84	HADFD01		HADFD10	HADFK11	HADFT44
	33	34	35	36		37	38	39

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					Lys-21 to Thr-26.	Thr-24 to Glu-29.																					Ser-90 to Lys-98.								
			-	2647	2648	2649	2650	2651																			2652								
				223 - 306	187 - 297	83 - 187	47 - 184																				1327 - 1698								
				20	51	52	53	54																			55								
				99066	741054	654831	826467	783049																			772423								
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<u>, , , , , , , , , , , , , , , , , , , </u>	Ser-90 to Lys-98.
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1, H0670: 1, S0044: 1, S3014: 1, L0439: 1, L0756: 1, L0779: 1, L0752: 1, L0757: 1, L0759: 1, H0444: 1, H0445: 1, H0595:	1, LO396: 1 and LO04: 1. H0521: 5, S0410: 3, S0476: 3, H0169: 3, L0766: 3, S0026: 3, H0542: 3, H0713: 2, T0049: 2, L0783: 2, L0773: 2, L0777: 2, L0783: 2, L0759: 2, L0588: 2, H0171: 1, H0158: 1, S0040: 1, H0295: 1, H0656: 1, H0341: 1, H0295: 1, H0656: 1, R0340: 1, S0132: 1, H0138: 1, S0007: 1, S0132: 1, H0138: 1, H0652: 1, H0673: 1, H0703: 1, H0591: 1, H0673: 1, H0708: 1, H0591: 1, H0673: 1, L0776: 1, L0657: 1, L0656: 1, S0374: 1, H0659: 1, L0659: 1, S0374: 1, H0659: 1, H0648: 1, H0672: 1, L0657: 1, L0659: 1, S0374: 1, H0539: 1, L0752: 1, S0031: 1, S0260: 1, H0444: 1, H0543: 1 and H0422: 1. S0358: 8, L0439: 5, H0052:	3, \$0360: 2, L0157: 2, L0438: 2, L0747: 2, L0438: 2, H0686: 1, T0049: 1, H0156: 1, \$0010: 1, H0581: 1, H0596: 1, H0038: 1, H0412: 1, \$0038: 1, H0412: 1, \$0038: 1, L0769: 1, L0775: 1, L0662: 1, L0766: 1, L0775:
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		1p36.23-p36.11														3q26.2-q27						
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			Met-1 to Trp-7,																			
	2657	2658	2659			2660						2661			2992	2663						
	35 - 142	205 - 258	106 - 282			187 - 246						285 - 383			128 - 325	358 - 414						
	09	61	62			63						2			65	99						
	823044	847117	561933			731929						844223			564230	823117						
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S0010: 1	AR060: 4, AR316: 3, AR089: 2 L0756: 9, L0740: 3, S0010: 1, S0051: 1, S6028: 1, L0770: 1, L0789: 1, L0438: 1 and L0600: 1.	S0010: 2, S0011: 2 and H0264: 1.	AR089: 17, AR316: 14, AR060: 12 L0777: 12, L0758: 9, L0439: 8, L0740: 7, L0779: 7, H0040: 6, L0748: 6, L0747: 6, H0038: 5, L0659: 5, L0595: 5, L0761: 4, L0776: 4, L0663: 4, L0565: 4, L0749: 4, L0663: 4, L0565: 4, L0749: 4, L0663: 4, L0565: 4, L0749: 4, L0665: 3, H0427: 3, H0050: 3, H0673: 3, H0068: 3, L0662: 3, L0596: 3, L0717: 2, H0057: 2, H0047: 2, L0771: 2, L0741: 2, H0641: 2, L0770: 2, L0749: 2, L0775: 2, L0805: 2, L0731: 2, H0641: 2, L0766: 2, L0731: 2, H0643: 2, L0765: 2, R0374: 2, H0643: 2, L0768: 1, H0685: 1, S0342: 1, R0343: 1, H0685: 1, S0342: 1, R0343: 1, H0649: 1, R0349: 1, H0341: 1, H0596: 1, L0118: 1, H0341: 1, H0596: 1, L0118: 1, H0588: 1, H079: 1, H0416: 1, H0588: 1, H0665: 1, R0366: 1, H0090: 1, H0606: 1, R039: 1, L0763: 1, H0699: 1, L0369: 1, L0763:
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59	09	61	62

	147781, 172471, 186580, 204200, 266600, 600760, 600760, 600761, 600761, 602594		
	16p12.1-q12.2		
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		Pro-57 to Phe-65.	
		2671	2672
	725 - 778	16 - 213	509 - 538
	73	74	75
	838174	837203	638587
	HAGB V25	HAGBV29	HAGCC87
	63	49	65

0010: 170041: L0809: Ind	p	, 1	10013: H0521: and	pı	5, H0032: , L0439: , L0591: 1, S0114: 1, S0114: 1, H0529: 1, L0782: 1, L0782: 1, H0144: 1, H0682: 1, L0779: 3, S0010: 1, H0542:	L0439: 1
AR060: 6 L0439: 4, L0519: 3, S0010: 2, L0731: 2, T0010: 1, T0041: 1, L0773: 1, L0766: 1, L0809: 1, H0682: 1, S0404: 1 and L0756: 1.	· ·	AR089: 5, AR316: 4, AR060: 4 S0010: 1 and S0152: 1	L0439: 2, L0759: 2, H0013: 1, S0010: 1, L0438: 1, H0521: 1, L0755: 1, L0731: 1 and L0758: 1.	S0010: 2, L0766: 2 and L0599: 2.	AR060: 6, AR316: 5, AR089: 3 L0794: 6, L0777: 3, H0032: 2, L0758: 2, L0438: 2, L0439: 2, L0758: 2, L0759: 2, L0591: 2, H0624: 1, H0717: 1, S0114: 1, H0661: 1, S0356: 1, S0010: 1, T0010: 1, S0040: 1, H0135: 1, L0763: 1, L0659: 1, L0764: 1, L0657: 1, L0659: 1, L0782: 1, H0659: 1, L0749: 1, H0144: 1, H0659: 1, L0749: 1, L0779: 1, L0754: 7, L0438: 2, S0010: 1, L0754: 1, L0780: 1, H0542: 1 and H0423: 1.	L0769: 3, L0438: 3, L0439:
		Asn-39 to Lys-44.				
	2673	2674	2675	2676	2678	2680
	150 - 275	31 - 231	42 - 161	91 - 168	744 - 836	45 - 140 110 - 121
	76	77	78	62	81	83
	561934	965095	751796	747697	724860	596804 566758
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	Gly-20 to His-33.			Asn-74 to Gln-79, Pro-89 to Val-104.	-											-						
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	2685													•	-																					
	198 - 305																																			
	88																																			
	589514																																			
	HAGDT30																																			
	78																																			

		124020, 134637, 157640, 174900, 236730, 278000, 278000, 600095, 600512, 601493
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	Thr-34 to Pro-39.	
	2686	2687
	90 - 209	223 - 240
	68	06
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	79	08

002001 47050	600511, 601556	103850, 114855, 116800, 140100, 140100, 192090, 192090, 192090, 192090, 245900, 245900, 276600,
+		16q22.1-q22.3
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	Met- 1 to Fne-6.	
	7088 7088 7088 7088 7088 7088 7088 7088	2689
	932 - 1051	165 - 260
	16	92
	793464	748222
	HAGEK37	HAGEK86
	81	82

600223						-			•										133780, 203100,	203100		154705, 190160,	771040,001134
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																		Ser-33 to Leu-38	Dr. 20 to Cue-30	FIO-20 to Cys-30.			
	2690	2691	2692	7/07														2603	2604	7024		2695	
	120 - 227	162 - 215	226 330	CCC - 077						-								175 336	10 113	18 - 113		959 - 909	
	93	94	30	5									-					70	2 2	16		86	
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	154705, 190160, 227646, 601154	
	3p24.3	
3, H0009: 3, L0769: 3, L0439: 3, H0441: 2, L0794: 2, L0741: 2, S0001: 1, S0045: 1, S6026: 1, H0565: 1, H0569: 1, H0504: 1, S0388: 1, H0399: 1, S0388: 1, L0767: 1, L0438: 1, H0520: 1, S0028: 1, L0438: 1, L04	AR060: 6, AR316: 5, AR089: 4 H0178: 4, L0351: 4, S0222: 3, H0009: 3, L0769: 3, L0439: 2, S0001: 1, S0045: 1, S6026: 1, S0010: 1, S0346: 1, H0194: 1, H0565: 1, H0569: 1, H0172: 1, S0388: 1, H0399: 1, S6028: 1, L0638: 1, L0767: 1, L0792: 1, L0438: 1, H0520: 1, S0028: 1 and L0595: 1.	AR053: 12, AR213: 10, AR290: 9, AR183: 9, AR270: 9, AR285: 9, AR271: 9, AR284: 8, AR241: 8, AR284: 8, AR268: 8, AR284: 8, AR268: 8, AR292: 7, AR184: 7, AR295: 7, AR175: 7, AR192: 7, AR310: 6, AR291: 6, AR243: 6, AR291: 6, AR243: 6, AR293: 6, AR282: 6, AR293: 6, AR309: 6, AR266: 6, AR309: 6, AR266: 6, AR309: 6, AR266: 6, AR309: 6, AR266: 5, AR296: 6, AR293: 5, AR296: 6, AR293: 5,
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	2696	2697
	959 - 909	13 - 198
	66	100
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5, AR232: 5, 5, AR089: 5,	5, AR206:	5, AR298:	5, AR104: 5, 5 AR240: 4	3, AR243.	4, AR219:					4, AR218:	4, AR227:	3, AR277:	3, AR258:	3, AR259:	3, AR248:	3, AR179:	2, AR231:		10, L0803: 4	4, L0747: 4, S0408: 3, L0770:	3, L0752: 3, H0445: 3, S0436:	3, S0276: 3, H0013: 2, L0471:	2, H0032: 2, S0422: 2, L0764:	2, L0517: 2, L0809: 2, L0665:	2, H0696: 2, H0436: 2, L0439:	2, L0750: 2, L0777: 2, S0308:	2, H0542: 2, H0717: 1, H0650:	1, H0662: 1, L0005: 1, S0356:	S0442: 1, S0444: 1, S0360:	S0300: 1, H0441: 1, H0587:	L0586: 1, S0010: 1, S0665:	H0581: 1, L0157: 1, S0388:	1, H0687: 1, H0328: 1, H0615:	1, H0124: 1, H0272: 1, H0412	T0004: 1, L0761: 1, L0667:	L0372: 1, L0646: 1, L0626:
AR096: AR299: AB367:	AR274:	AR294:	AR316:	AR186:	AR253:	AR055:	AR289:	AR244:	AR247:	AR185:	AR240:	AR237:	AR204:	AR300:	AR234:	AR256:	AR233:	AR061:	99L0T	4, L074	3, L075	3, S027	2, H003	2, L051	2, H069	2, L075	2, H054	1, H066	1, S044	1, S030	1, L058	1, H058	1, H068	1, H012	1, T000	11, LU3/
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		Cys-26 to Asp-31.		Gln-47 to Gly-52. Asn-34 to Gly-39.		
	2698	2699	2700	2701	2703	2704
	266 - 283	208 - 486	243 - 278	830 - 1192	183 - 287	342 - 359
	101	102	103	104	901	107
	821316	861680	604536	780112	634611	838059
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	91	92	93	94	96	97

			143890, 151440, 600173, 600276, 600310, 600310, 601604, 601843
			19p13.1
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		7 7	
	2705		2706
	28 - 126		59 - 130
	108		109
	655435		845963
	HAGHR18		HAGIB90
	86		66

, H0510: 2, F , L0805: 2, L , L0805: 2, L , L0605: 2, S , H0171: 1, 1 , H0171: 1, 1 , H052: 1, 1 , H053: 1, 1 , L0764: 1, L , L0806: 1, L , L0790:	AR060: 9, AR316: 7, AR089: 5 H0599: 1	H0599: 5, L0777: 2, L0485: 2, H0097: 1, H0706: 1, L0774: 1, L0666: 1, L0747: 1, L0780: 1 and L0759: 1.	L0731: 12, H0545: 10, H0341: 7, L0769: 7, L0748: 7, L0751: 7, L0750: 7, S0144: 6, L0766: 6, L0747: 6, L0752: 6, L0758: 6, H0295: 5, S0360: 5, S0344: 5, L0775: 5, L0744: 5, L0754: 5, L0755: 5, H0657: 4, H0305: 4, S0278: 4, H0424: 4, H0213: 4, L0774: 4, L0659: 4, L0749: 4, S0007: 3, S0045: 3, S0046: 3, L0622: 3, H0266: 3, H0546: 3, L0622: 3, H0677: 3, H0059: 3, L0803: 3, S3014: 3, L0777: 3, L0757: 3, H0171: 2,
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	2707	2708	2709
	185 - 304	295 - 600	85 - 234
·	110	111	112
	835871	848816	847014
	HAHEM51	HAHSA76	HAHSD51
	100	101	102

			S0040: 2, H0663: 2, S0358: 2,	
			S0408: 2, S0132: 2, L0717: 2,	
			H0351: 2, S0222: 2, H0333: 2,	
			H0486: 2, H0101: 2, T0048: 2,	
			H0318: 2, H0052: 2, H0041: 2,	
			H0009: 2, L0157: 2, H0510: 2,	
			H0622: 2, H0604: 2, H0644: 2,	
			H0617: 2, H0040: 2, H0494: 2,	
			S0438: 2, H0633: 2, L0770: 2,	
			L0761: 2, L0764: 2, L0521: 2,	
			L0375: 2, L0776: 2, L0783: 2,	•
			S0126: 2, H0670: 2, S0152: 2,	
	•		S0027: 2, S0032: 2, L0753: 2,	
			1 0759 2 H0624: 1 H0170: 1	
			H0204: 1 H0650: 1 H0381: 1	
			110274: 1, 110030: 1, 110301: 1,	
			S0116: 1, H0484: 1, H0233: 1,	
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			H0108: 1, T0082: 1, S0010: 1,	
			S0049: 1, H0173: 1, H0196: 1,	
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			H0649: 1, S0142: 1, S0422: 1,	
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			L0644: 1, L0771: 1, L0773: 1,	
			, L0653: 1,	
			L0655: 1, L0658: 1, L0809: 1,	
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			H0520: 1, H0690: 1, H0683: 1,	
			H0659: 1, S0328: 1, S0378: 1,	
			H0134: 1, S0404: 1, H0436: 1,	

	17pter-p13.1			2
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1 I S 1	Ser-33 to Ala-45, 32 Glu-48 to Lys-58. 2 2 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		Pro-19 to Ser-28.	
	2710	2711	2712	2713
	137 - 355	286 - 393	90 - 194	363 - 380
	113	114	115	116
	874298	741086	852223	834523
	HAIBR76	HAIBT20	HAIBV91	HAICE62
	103	104	105	106

		176000, 261640, 602574, 602574
		11q23.3
S0212: 1, S0442: 1, S0358: 1, S0132: 1, S0476: 1, S0278: 1, L0021: 1, H0599: 1, H0036: 1, H0123: 1, H0559: 1, H0559: 1, H0036: 1, H0123: 1, H00373: 1, H0031: 1, H0031: 1, H0032: 1, H0211: 1, H0090: 1, H0040: 1, H0242: 1, L0742: 1, L0742: 1, L0742: 1, L0743: 1, L0743: 1, L0743: 1, L0744: 1, L0744: 1, L0744: 1, L0744: 1, L0744: 1, L0746: 1, L0746: 1, L0748: 1, L0740: 1, L0748: 1, L0740: 1, L0748: 1, L0740: 1, L0753: 1, L0553: 1, L0553: 1, H0673: 1, L0593: 1, H0672: 1, L0593: 1, H0672: 1, L0593: 1, H0672: 1, L0748: 1, L0740: 1, L0750: 1, L0755: 1, L0593: 1, H0672: 1, H0673: 1, H0673	L0754: 4, S0132: 3, S0444: 2, L0750: 2, H0689: 1, H0522: 1 and H0423: 1.	L0740: 9, L0439: 7, S0360: 5, L0770: 4, L0665: 4, L0748: 4, L0747: 4, L0752: 4, H0431: 3, H0574: 3, L0105: 3, H0553: 3, L0766: 3, L0774: 3, L0105: 3, L0765: 3, L0774: 3, L0774: 3, L0774: 3, L0774: 3, L0774: 3, L0731: 3, H0662: 2, S0007: 2, S0046: 2, L0717: 2, H0560: 2, H0549: 2, L0754: 2, L0756: 2, L0665: 2, L0665: 2, L0754: 2, L0756: 2, L0605: 2, L0362: 2, S0242: 2, H0686: 1, S0212: 1, S0420: 1, R0358: 1, R0376: 1, L0021: 1, H0497: 1, H0497: 1, H0421: 1, H0596: 1, H0531: 1, H0421: 1, H0596: 1, H0531: 1, H0521: 1, H0590: 1, H0590: 1, H0590: 1, H0590: 1, H0590: 1, H0591: 1, H0590:
881111881111811111111111111111111111111		
	2714	2715
	140 - 259	232 - 312
	117	118
	637491	772534
	HAICL90	HAICV44
	107	108

		143890, 151440,
		19p13.1
1, L0041: 1, H0086: 1, L0471: 1, S6028: 1, H0687: 1, S0003: 1, H0039: 1, L0194: 1, H0032: 1, H0124: 1, S0036: 1, H0124: 1, S0036: 1, H0124: 1, H0124: 1, H0059: 1, H0494: 1, H0646: 1, S0208: 1, L0763: 1, L0769: 1, L0764: 1, L0374: 1, L0389: 1, L0775: 1, L0374: 1, L0389: 1, L0775: 1, L0352: 1, H0520: 1, H0658: 1, H0672: 1, H0539: 1, L0602: 1, H0696: 1, H0555: 1, S0206: 1, L0744: 1, L0753: 1, L0759: 1, L0744: 1, L0753: 1, L0759: 1, L0589: 1, L0366: 1, S0196: 1, S0194: 1, S0196: 1, and S0424: 1.	L0439: 8, L0777: 8, L0666: 4, S0010: 3, H0615: 3, L0637: 3, L0659: 3, L0664: 3, H0648: 3, L0748: 3, L0731: 3, S0132: 2, S0476: 2, S0222: 2, L0665: 2, H0497: 2, L0774: 2, L0655: 2, H0144: 2, L0738: 2, L0754: 2, L0757: 2, L0758: 2, S0194: 2, H0171: 1, H0269: 1, H0570: 1, H0569: 1, H0550: 1, H0569: 1, H0628: 1, H0427: 1, H0569: 1, H0628: 1, H0427: 1, H0563: 1, H0628: 1, H0628: 1, L0805: 1, L0769: 1, L0753: 1, L0755: 1, L0769: 1, L0753: 1, L0755: 1, L0769: 1, S0031:	L0776: 6, L0775: 5, L0769:
	4 6 6 6 6 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	Trp-23 to Ala-28,
	2716	2717
	255 - 332	126 - 404
	116	120
	847015	780114
	HAIDP45	HAJAB88
	109	110

600173, 600276, 600310, 600310, 601604, 601843			21q22.2 176261, 601399
4, L0779; 4, S0358; 3, H0038; 3, L0809; 3, L0747; 3, H0445; 3, H0620; 2, L0763; 2, L0766; 2, L0766; 2, L0766; 2, L0766; 2, L0766; 2, L0755; 2, L0770; 2, L0777; 2, L0766; 2, L0755; 2, L0777; 2, L0755; 2, L0759; 2, H0171; 1, H0556; 1, H0688; 1, H0125; 1, H0661; 1, H0638; 1, H0125; 1, H0392; 1, H0013; 1, H0599; 1, S0010; 1, H0318; 1, S0049; 1, H0012; 1, H0771; 1, H0551; 1, H0012; 1, L0771; 1, H0551; 1, L0806; 1, L0647; 1, L0806; 1, L0647; 1, H0660; 1, L0782; 1, L0647; 1, H0660; 1, H0666; 1, S0126; 1, H0543; 1 and H0422; 1, L0757; 1, L0757; 1, L0757; 1, L0757; 1, L0745; 1, L0757; 1, L0745; 1, L0757; 1, L0475; 1, L0745; 1, L0757; 1, L0485; 1, H0543; 1 and H0422; 1.	L0770: 4, H0457: 3, L0756: 2, L0759: 2, H0663: 1, H0587: 1, H0013: 1, H0427: 1, H0318: 1, H0594: 1, S0003: 1, H0561: 1, H0529: 1, L0805: 1, L0776: 1, L0655: 1, L0757: 1, L0758: 1, L0755: 1, L0757: 1, L0758: 1 and L0594: 1.	L0766: 4, H0657: 1, H0486: 1, H0013: 1, H0644: 1, L0456: 1, H0560: 1, L0662: 1, L0804: 1, L0774: 1, L0793: 1, S0328: 1, L0439: 1, L0749: 1 and H0543: 1.	L0777: 5, L0766: 3, L0803: 21c 3, L0439: 3, S0360: 2, L0598: 2, L0666: 2, L0748: 2, T0049:
Pro-39 to Gly-65.	·	Val-34 to Glu-39.	Thr-22 to Lys-30.
	2718	2719	2720
	300 - 407	282 - 500	90 - 203
	121	122	123
	716168	588488	610051
	HAJAZ56	HAMFC67	HAMFQ38
	Ξ	112	113

	107680, 107680, 107680, 107680, 107680, 107680, 107720, 133780, 147791, 159555, 168000, 186740, 18625, 203750, 261640, 600048, 601382, 602574, 602574	
1, \$0134: 1, \$0116: 1, L0717: 1, H0586: 1, H0486: 1, H0575: 1, H0510: 1, H0553: 1, H0560: 1, \$0422: 1, L0763: 1, L0769: 1, L0521: 1, L0767: 1, L0768: 1, L0775: 1, L0663: 1, \$0374: 1, L0438: 1, H0520: 1, H0682: 1, \$0328: 1, L0740: 1, L0757: 1, \$0192: 1 and H0543: 1.	AR060: 2, AR316: 2, AR089: 2 L0766: 5, S0418: 3, H0144: 3, L0777: 3, L0758: 3, L0769: 2, L0771: 2, L0378: 2, L0663: 2, L0771: 2, L0378: 2, L0663: 1, S0134: 1, H0657: 1, H0484: 1, S0420: 1, L0717: 1, S0222: 1, H0270: 1, L0717: 1, S0222: 1, H0263: 1, H0266: 1, H0687: 1, H0252: 1, H0604: 1, H0628: 1, H0252: 1, H0606: 1, H0135: 1, H0617: 1, H0606: 1, H0561: 1, L079: 1, L076: 1, L0794: 1, L0774: 1, L0776: 1, L0789: 1, L0774: 1, L0776: 1, L0789: 1, L0774: 1, L0776: 1, L0438: 1, L0774: 1, L0776: 1, L0789: 1, L0774: 1, L0779: 1, L0751: 1, L0777: 1, L0751: 1, H0596: 1, L0777: 1, L0751: 1, H0596: 1, L0777: 1, L0779: 1, L0751: 1, L0777: 1, L0779: 1, L0752:	AR089: 15, AR316: 11, AR060: 9 L0540: 5, S0002: 2, L0663: 2, L0596: 2, S0354: 1, S0376: 1, S0360: 1, H0574: 1, S0049: 1, L0738: 1, L0041: 1, H0051: 1, S0318: 1, S0316: 1, S0036: 1, S0426: 1, L0764: 1, L0662:
1, S0134: 1 1, H0586: 1, H0510: 1, S0422: 1 1, L0521: 1, L0775: 1, L0438: 1, S0328: 1	AR060: 2, AR AR089: 2 L0766: 5, S04 3, L0777: 3, L0 2, L0771: 2, L0 2, L0740: 2, L0 1, S0134: 1, H0 1, S0420: 1, L0 1, H0270: 1, H0 1, H0252: 1, H0 1, H0100: 1, H0 1, L0796: 1, L0 1, L0796: 1, L0 1, L0796: 1, L0 1, L0793: 1, L0 1, L0791: 1, L0 1, L0791: 1, S0328: 1, L0 1, L0791: 1, S0338: 1, L0 1, L0791:	
	2721	399 2722 Ala-17 to Gln-23
	783864 124 644 - 697	664478 125 199 - 399
	114 HAMGG01 783	115 HANGB24 664

				108725, 120700, 133171, 136836, 143890, 145981, 147141, 147670,
				19p13.3-p13.2
1, L0794: 1, L0804: 1, L0774: 1, L0776: 1, L0666: 1, S0374: 1 and L0748: 1.	AR089: 33, AR316: 23, AR060: 13 L0109: 1, S0318: 1, S0316: 1, H0163: 1, L0756: 1 and L0601: 1.	H0042: 1	L0766: 8, H0677: 6, L0759: 5, L0794: 4, L0749: 4, L0662: 3, L0659: 3, L0809: 3, L0787: 5, H0436: 3, L0747: 3, H0716: 2, S0358: 2, H0486: 2, H0402: 2, L0769: 2, L0764: 2, L0776: 2, L0776: 2, L0776: 2, L0776: 2, L0776: 2, L0776: 2, L0778: 2, L0778: 2, L0738: 1, H0501: 1, S0222: 1, H0502: 1, H0501: 1, S0222: 1, H0502: 1, H0650: 1, T0114: 1, H0013: 1, T0048: 1, H0052: 1, L0664: 1, L0761: 1, L0775: 1, L0768: 1, L0761: 1, L0775: 1, L0768: 1, L0768: 1, L0768: 1, L0778: 1, L0768: 1, L0778: 1, L0664: 1, L0666: 1, L0664: 1, L0666: 1, H0519: 1, H0689: 1, H0435: 1, H0519: 1, L0750: 1, H0518: 1, L0739: 1, L0750: 1, L0748: 1, H0518: 1, L0439: 1, L0750: 1, S0434: 1, And L0599: 1, S0434: 1, And L0599: 1, S0434: 1, And L0599: 1, L0750: 1, L0759: 1, L075	AR089: 58, AR316: 46, AR060: 36 L0439: 18, H0521: 9, H0618: 6, H0253: 6, H0457: 5, L0438:
	Lys-18 to Gly-23, Arg-25 to Ser-31.			
	2723	2724	2725	2726
	198 - 317	50 - 154	125 - 283	247 - 384
	126	127	128	129
	847018	840584	926609	762803
	HANKC93	HAPAD35	HAPBR13	HAPBU09
	116	117	118	611

14/6/U, 14/6/U, 15/1440, 164953, 188070, 231670, 600276, 600957, 601238, 601846, 602216, 602477	142983, 222300	142983, 222300	156845, 156845,
	4p16.1	4p16.1	3p13-q13.33
4, L0743: 4, L0748: 4, H0341: 3, H0255: 3, S0222: 3, L0794: 3, L0741: 2, S0408: 2, S0444: 2, S0402: 2, L0437: 2, L0769: 2, L0769: 2, L0763: 2, L0833: 2, L0833: 2, L0533: 2, L0749: 2, S0031: 2, L0591: 2, L0749: 2, S0031: 2, L0591: 2, L0749: 2, S0031: 2, L0591: 2, H0625: 1, H0717: 1, R0340: 1, H0625: 1, H0033: 1, H0052: 1, H0068: 1, H0069: 1, H0063: 1, H0668: 1, H0668: 1, H0668: 1, L0779: 1, L0766: 1, L0766: 1, L0778: 1, L0659: 1, L0779: 1, L0777: 1, L0744: 1, L0779: 1, L07	L0752: 7, L0439: 5, L0438: 2, S0356: 1, H0013: 1, H0042: 1, T0023: 1, H0166: 1, H0591: 1, L0803: 1, L0515: 1, L0659: 1, L0647: 1, L0789: 1, H0539: 1, L0731: 1 and H0352: 1.	L0752: 7, L0439: 5, L0438: 2, S0356: 1, H0013: 1, H0042: 1, T0023: 1, H0166: 1, H0591: 1, L0803: 1, L0515: 1, L0659: 1, L0647: 1, L0789: 1, H0539: 1, L0731: 1 and H0352: 1.	AR277: 18, AR244: 15,
	Ser-20 to Asp-25.	Ser-20 to Asp-25.	Leu-36 to Ser-49.
	2727	2728	2729
	403 - 510	403 - 510	1872 - 2084
	130	131	132
	789544	864890	835554
	HAPBU86	HAPBU86	HAPNJ33
	120	121	122

		AR269: 13, AR275: 13, AR183: 13, AR232: 13, AP246: 12, AP238: 12	600151
		13, AR2/3: 13, AR232: 12, AR238:	161000
		13, AR232:	_
		12 AB238.	
		12, AN230.	
		AR227: 12, AR234: 12,	
		11, AR218:	
		11, AR243:	
		10, AR186:	
		AR268: 10, AR061: 10,	
		AR290: 10, AR226: 10,	
		10, AR309:	
		10, AR053:	
	•	9. AR299:	
		9, AR312:	
		8. AR233:	
		8 AR198:	
		8 AR185:	
		8 AD187.	
		8, AR182.	
		7, AR267	
		7. AR249:	
		7, AR280:	
		7. AR177:	
		6, AR241:	
		6, AR213:	
		AR096: 6, AR205: 6,	
		AR263: 6, AR240: 6,	
		AR291: 6, AR175: 6,	
	-	AR253: 5, AR033: 5,	
		AR295: 5, AR265: 5,	
		AR104: 5, AR286: 5,	
		4, AR285:	
		4, AR281:	
		AR283: 3, AR294: 3,	

AR258: 3, AR256: 3, AR259: 3, AR179: 3 S0358: 11, H0545: 8, H0622: 8, H0551: 7, H0547: 7, H0575: 6, L0666: 6, L0664: 6, S0436:	6, H0542: 6, H0584: 5, L0769: 5, L0662: 5, L0766: 5, L0789: 5, L0438: 5, H0521: 5, S0406: 5, L0756: 5, L0777: 5, L0755: 5, H0543: 5, H069: 4, H0251: 4, H0546: 4, H0286: 4, H0617:	4, L0770: 4, L0638: 4, L0/61: 4, L0657: 4, L0659: 4, L0663: 4, L0439: 4, L0740: 4, L0751: 4, H0665: 4, H0599: 3, H0544: 3, H0355: 3, H0412: 3, H0100: 3, L0646: 3, L0771: 3, L0724: 1, L0725: 3, L0721: 3, L0724: 1, L0725: 3, L0726: 3, L07	3, L5623: 3, L0565: 3, H0519: 3, L0747: 3, L0750: 3, L0752: 3, L0747: 3, L0758: 3, S0434: 3, S0424: 3, H0265: 2, H0556: 2, H0341: 2, H0458: 2, S0420: 2, S0360: 2, S0410: 2, H0618: 2, H0613: 2, H0618: 2, H06	2, H0051: 2, H0083: 2, H0375: 2, S0250: 2, H0039: 2, H0030: 2, H0623: 2, H0509: 2, H0413: 2, H0623: 2, S0426: 2, L0763: 2, L5565: 2, L0772: 2, L0768: 2, L0774: 2, L0773: 2, L0768: 2, L0774: 2, L0551: 2, L0806: 2, L0553: 2, L0551: 2, L05	2, L0651: 2, L0680: 2, L0650: 2, L0665: 2, H0435: 2, H0436: 2, H0539: 2, L0597: 2, L0597: 2, L0591: 2, L0593: 2, L0361: 2, H0423:
AR258: AR259: S0358: 8, H055 6, L066	6, HO 5, LO 5, LO 5, LO 5, HO 6, HO	7, 7, 7, 7, 7, 7, 7, 7, 7, 7, 7, 7, 7, 7	7, 12 3, 12 3, 10 3, 10 2, 10 2, 10 2, 10 2, 10 2, 10 2, 10 3, 10 4, 10 5, 10	2, 10 2, 10 2, 10 2, 10 1, 1	2, LC 2, LC 2, SO 2, SO 2, LC 2, LC 2, LC

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170:	713:	384:	300:	154:	045:	l, H0392:	ı, L0622:	l, H0427:	048:	052:	046	3012:	010:	179:	338:	615:)628:	124:	112:	1, H0494:	, H0130:	, H0529:	L4747:	L5566:	L0374:	381:	, L0540:	374:	:0690	.0990	555:	2612:	028:	753:	0445:	2990	1412:	
4: 1, H0	J: 1, HU 6: 1 TO	2: 1, HO	2: 1, H0	5: 1, S03	3: 1, S0	1: 1, HC	3:1, L0	0: 1, HC	S0010: 1, T0048:	, S0049: 1, H0052:	7: 1, HC	0: 1, HC	1, H0373: 1, T0010:	1, H0267: 1, H0179	1, S0336: 1, S0338:	2:1.H0	4: 1, H(5. 1 HC	58: 1. SC	2: 1, H0	8: 1, HC		•		1:1, L0	1, L0767: 1, L0381:	6: 1, L0	H0144: 1, S0374:	39: 1, H(58: 1, H	8: 1, H0	26: 1, L(4: 1, SO	9: 1, L0	14: 1, H(53: 1, H(22: 1, SC	
2, H062	1, SUU4(1, 11005 1, S0282	1, H040	1, \$0356	1, H073	1, H0441: 1	1, H0333: 1	1, H0250: 1	1, S001	1, S004	1, H059	1, H005	1, H037	1, H026	1, S033	1. H025	1. H064	1 1.045	1, <u>200</u>	1, T0042: 1	1, S043	1, H0538: 1	1, L0371: 1	1, L3905: 1	L0667: 1, L0641: 1,	1, L076	•	1, H014	1, H068	1, H06;	1, S026	1, H062	1, \$301	1, L074	1, H044	1, H065	1, H042	56: 1.
2, H0506: 2, H0624: 1, H0170:	1, H0685: 1, S0040: 1, H0/15; 1, H0687: 1, H0686: 1, 10428	1, F0037. 1, 110030. 1, E0423. 1, S0212: 1, S0282: 1, H0384:	1, H0663: 1, H0402: 1, H0300	1, S0418: 1, S0356: 1, S0354:	1, H0728: 1, H0733: 1, S0045:	1, H0549:	H0600:	H0485:	1, H0098:	H0505:	H0194: 1, H0597: 1, H0046:	H0081: 1, H0050: 1, H0012	1 H0024:	H0266:	H0188:	S0340: 1, H0252: 1, H0615:	H0213: 1, H0644: 1, H0628:	H0606: 1 1 0455: 1 H0124:	H0063: 1, H0268: 1, S0112:	L0351: 1.	H0625: 1, S0438: 1	S0344:	L0369:	L0637: 1	L0667:	L0648:	L0378:	L0543: 1	H0593: 1, H0689: 1, H0690:	H0659: 1, H0658: 1, H0660:	S0152: 1, S0268: 1, H0555:	1, H0478: 1, H0626: 1, L0612	1, S0432: 1, S3014: 1, S0028:	1, L0748: 1, L0749: 1, L0753:	1, L0759: 1, H0444: 1, H0445	1, L0592: 1, H0653: 1, H0667	1, S0192: 1, H0422: 1, S0412:	and S0456: 1
2,1	<u>-</u> -	<u> </u>	<u></u>	1,5		<u></u>				<u> </u>	<u> </u>	<u> </u>	<u> </u>	-	<u></u>	· —		_	<u>-</u>		1	1,		<u> </u>	<u>1</u> ,	<u>-</u>	<u>1</u> ,	<u>–</u>	<u></u>	<u>1</u> ,	1,	<u>-</u>		1,	1,	Ή,	<u>-</u>	1
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11pter-p15.5	
AR060: 4, AR316: 4, AR089: 4 L0717: 5, L0794: 4, L0803: 3, L0770: 3, L0766: 3, L0779: 2, L0764: 2, L0761: 2, L0764: 2, L0769: 2, L0777: 2, L0769: 2, L0777: 2, L0769: 2, L0769: 1, H0677: 1, H0677: 1, H0677: 1, H0639: 1, H0677: 1, H0639: 1, H0024: 1, H0171: 1, H0173: 1, H0639: 1, H0020: 1, L0667: 1, L0749:	AR060: 6, AR089: 3, AR316: 3, AR215: 3, AR311: 3, AR213: 3, AR223: 3, AR282: 3, AR207: 3, AR197: 2, AR296: 2, AR254: 2, AR168: 2, AR266: 2, AR270: 2, AR216: 1, AR283: 1, AR216: 1,
Glu-45 to Thr-63.	
2730	2731
701 - 1009	205 - 315
133	134
790340	834384
HAPNL62	HAPNO50
123	124

	602134
	2p24.3-p24.1
ARI71: 1, AR165: 1, AR096: 1, AR264: 1 L0439: 10, L0779: 6, L0794: 5, L0774: 4, L0375: 4, L0747: 3, L0803: 3, L0809: 3, L0771: 3, L0809: 3, L0771: 3, L0809: 3, L0771: 3, L0809: 2, H0652: 2, H0652: 2, H0652: 2, L0768: 2, L0766: 2, H0672: 2, L0768: 2, L0766: 2, H0672: 2, L0743: 2, H0624: 1, H0294: 1, H0294: 1, H0294: 1, H0294: 1, H0294: 1, H0498: 1, L0005: 1, H0733: 1, H0438: 1, L0005: 1, H0675: 1, S0010: 1, S0049: 1, H0675: 1, H0687: 1, H0620: 1, S0049: 1, H0644: 1, L0434: 1, H0551: 1, H04413: 1, L0434: 1, L0775: 1, L0769: 1, L0776: 1, L0776	AR089: 25, AR316: 20, AR060: 16 L0766: 13, L0770: 9, S0003: 8, L0439: 8, L0754: 8, L0665: 7, L0731: 6, L0485: 6, L0776: 5, L0740: 5, H0641: 4, L0771: 4, H0521: 4, L0741: 4, L0752: 4, S0356: 3, S0360: 3, L0717: 3, H0013: 3, H0014: 3, L0598: 13, L0749: 3, L0777: 3, L0749:
	2732
	33 - 167
	135
	702037
·	HAPNY10
	125

	•	
2, L0005: 2, H0428: 2, S0408: 2, S0408: 2, S0408: 2, L0369: 2, L0662: 2, L0659: 3, L0604: 1, H0556: 1, H0574: 1, H0574: 1, H0674: 1, H0676: 1, H0678: 1, H0650:	, L0608:	H0575: , S0192:
S0424: 3, S0212: 2, L0005: S0358: 2, S0376: 2, S0408: S0410: 2, S0222: 2, H0575: H0421: 2, H0641: 2, H0642: 2, L0599: L0772: 2, L0773: 2, L0659: L0772: 2, L0750: 2, L0599: 3, H0661: 1, H0638: 1, S0218: H0661: 1, H0638: 1, S0218: H0641: 1, H0638: 1, S0218: H0641: 1, H0638: 1, S0418: H0641: 1, H0638: 1, S0418: H0641: 1, H0638: 1, H0574: H0448: 1, H0655: 1, H0674: H0690: 1, H0625: 1, H0674: H0590: 1, H0625: 1, L0640: L0768: 1, L0669: 1, H0625: 1, L0640: L0784: 1, L0803: 1, L0774: L0784: 1, L0803: 1, L0774: L0784: 1, L0803: 1, L0774: L0784: 1, L0803: 1, L0793: L0519: 1, H0660: 1, H0660: 1, H0660: 1, H0660: 1, H0660: 1, H0650: 1, H0650: 1, L0783: 1, L0783: 1, L0783: 1, L0783: 1, H0650: 1, H0660: 1, H0660: 1, H0650: 1, L0755: 1, L0781: 1, L0782: 1, L0783: 1, L0782: 1, L0783: 1, L0783: 1, L0783: 1, L0783: 1, L0783: 1, H0650: 1, H0660: 1, L0783: 1, L07	L0599: 1 S0011: 1	L0748: 4, , L0665: 2
S0424: 3, S0358: 2, S0358: 2, S0410: 2, H0421: 2, H0421: 2, L0772: 2, L0772: 2, L0595: 2, L0758: 2, L0758: 2, L0758: 2, L0758: 1, H0661: 1, H0441: 1, H0441: 1, H0486: 1, H0880:	1, L0592: 1, L0599: 1, L0608: 1, L0601: 1, S0011: 1 and H0653: 1.	H0250: 7, L0748: 4, H0575: 2, H0457: 2, L0665: 2, S0192:
<u>κ, γ, /u>	` ', '	, , T
		2733
		86 - 9
		99
		136
		847020
		HAPPW83
		126

			4p15.3-p15.1
2, H0624: 1, H0295: 1, S0442: 1, H0580: 1, H0609: 1, H0050: 1, S0022: 1, H0628: 1, H0550: 1, H0090: 1, H0090: 1, L0642: 1, L0662: 1, L0767: 1, L0665: 1, L0669: 1, L0542: 1, L0666: 1, S0374: 1, H0547: 1, H0521: 1, H0521: 1, H0521: 1, H0521: 1, H0522: 1, L0740: 1 and S0026: 1.	5, AR316: 4 4 59, L0745: 9, 5: 3, H0547: 3 8: 2, L0646: 2 9: 2, L0362: 2 9: 2, L0362: 2 9: 1, H0661: 1 0: 1, H0546: 1 1, H055: 0: 1, H055: 0: 1, H0560: 1 8: 1, L0762: 1 8: 1, L0762: 1 8: 1, L0762: 1 8: 1, L0762: 1 9: 1, L0866: 1 9: 1, L0866: 1 9: 1, L0866: 1 9: 1, L0866: 1 9: 1, L0779: 1 9: 1, L0589: 1	1	40: 17, AR243: 15, 46: 14, AR271: 14, 42: 12, AR272: 11, 97: 7, AR161: 7, 62: 7, AR245: 7, 63: 6, AR282: 6,
2, HG 1, HG 1, LO 1, LO 1, LO 1, LO 1, HG 1, HG	AR089: AR060: L0754: 5, L0666: 2, S0438: 2, L0649: 1, R013: 1, H059: 1, H059: 1, H059: 1, L015 1, L059: 1, L079: 1, L079	AR060: AR089: H0575	AR240: AR246: AR242: AR197: AR162: AR163:
			Pro-28 to Ser-35.
	2734	2735	2736
	286 - 336	300 - 338	1324 - 1485
	137	138	139
	280802	637494	752580
	HAPQJ73	НАРQК26	HAPQU71
	127	128	129

													-																							·			
6. AR053:	5, AR195:							3, AR096:	3, AR264:	3. AR312:	3. AR250:	3 AR316:	2, AR291:		2, AR174:	2 AR055:		2, AR181		2, AR213:			_		_	_	Π	_	_	52: 1, AR227: 1	H0575: 7, L0751: 5, H0042:	4, L0664: 3, L0665: 3, H0046:	2, H0024: 2, H0658: 2, L0599:	2, H0685: 1, H0663: 1, H0477:	1, S0016: 1, L0639: 1, L0662:	1, L0775: 1, L0806: 1, L0527:	1, L0657: 1, L0666: 1, H0690:	1, H0670: 1, H0666: 1, H0672:	I and LU/55: 1.
JAR198:	AR165:	AR309:	AR204:	AR164:	AR274:	AR254:	AR201:	AR060:	AR311:	AR235:	AR039:	AR263:	AR205:	AB104:	AR247	AR267	AR217	AR300:	AR185	AR187:	AR283:	AR308:	AR212:	AR261:	AR228:	AR222:	AR199:	AR224:	AR233:	AR252:	OH H05	4, L0	2, HC	2, HC	1, S0	1, LC	1, LC	1, H(1 am
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4p15.3-p15.1																											-												
		12, AK2/2: 11,	/, AKIOI: /,		6, AK282: 6,				5, AR166: 5,	4, AR193: 4,	4, AR313: 4,	4, AR275: 4,	4, AR207: 3,	3, AR096: 3,	3, AR264: 3,	3, AR312: 3,					2, AR174: 2,	2, AR055:	2, AR229:		2, AR176: 2,	2, AR213:	(1	-	_		_	1, AR171: 1,	1, AR289: 1,	_		_	H0575: 7, L0751: 5, H0042:	4, L0664: 3, L0665: 3, H0046:	2, H0024: 2, H0658: 2, L0599:
AR240:	AR246:	AR242:	AR19/:	AK162:	AK165:	AK196:	AR165:	AR309:	AR204:	AR164:	AR274:	AR254:	AR201:	AR060:	AR311:	AR235:	AR039:	AR263:	AR205:	AR104:	AR247:	AR267:	AR217:	AR300:	AR185:	AR182:	AR283:	AR308:	AR212:	AR261:	AR228:	AR222:	AR199:	AR224:	AR233:	AR252:	H0575	4, L066	2, H002
Pro-28 to Ser-35.																																							
2737																																							
1324 - 1485																																							
140																																							
864781																																							
HAPQU71	,																																						
130																																							

2, H0685: 1, H0663: 1, H0477: 1, S0016: 1, L0639: 1, L0662: 1, L0775: 1, L0806: 1, L0527: 1, L0657: 1, L0666: 1, H0690: 1, H0670: 1, H0666: 1, H0672: 1 and L0755: 1. L0794: 2, L0789: 2, L0663: 2, L0779: 2, H0575: 1, L0761: 1, L0809: 1, L0805: 1, L0748: 1, and L0731: 1.	H0305: 4, L0805: 3, L0747: 3, H0040: 2, L0803: 2, S0410: 1, H0580: 1, S0300: 1, H0587: 1, H0486: 1, H0575: 1, H0318: 1, S6028: 1, H0598: 1, H0551: 1, H0560: 1, L0773: 1, H0435: 1, H0539: 1, L0758: 1, S0436: 1, H0542: 1 and H0543: 1.	AR089: 26, AR316: 21, AR060: 16 L0439: 7, L0809: 4, L0751: 4, L0659: 3, H0144: 3, L0438: 3, L0758: 3, S0360: 2, L0766: 2, L0803: 2, H0658: 2, L0743: 2, L074: 2, L0754: 2, L0777: 2, H0716: 1, H0657: 1, H0341: 1, S0212: 1, H0664: 1, H0125: 1, S0376: 1, H0570: 1, H0639: 1, H0623: 1, H0596: 1, T0010: 1, H0510: 1, R0596: 1, H0039: 1, H0622: 1, H0553: 1, L0655: 1, H0090: 1, L0772: 1, L0649: 1, L0764: 1, L0772: 1, L0649: 1, L0764: 1, L0662: 1, L0775: 1, L0805: 1, L0776: 1, L0657: 1, H0648:
2, H 1, S 1, L 1, L 1, H 1, H 1, H 1, L 1, L 1, L 1, L 1, L	3, H 1, 1 1, 1 1, 1 1, 1	ARA ARA 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
2738	2739	2741
20 - 133	515 - 568	172 - 288 251 - 421
141	142	143
610052	779053	847021
HAPQW18	НАРОХ44	HAPSH37
131	132	133

		118800, 123660, 125660, 125660, 193500, 193500, 193500, 193500, 201460, 205100, 237300, 262000, 600266, 601277
	×	2435
1, H0539: 1, H0522: 1, H0696: 1, H0436: 1, L0748: 1, L0749: 1, L0748: 1, H0543: 1 and H0423: 1. H0595: 2, H0392: 1, H0587: 1, H0587: 1, H0587: 1, H0587: 1, H0587: 1, H0589: 1, H0599: 1, L0759: 1, L0759: 1, L0759: 1, L0754: 1, L0747: 1, S0390: 1, L0754: 1, L0747: 1, S0390: 1, L0754: 1, L0747: 1, L0747: 1, L0747: 1, L0747: 1, L0748: 1, L0747: 1, L0747: 1, L0747: 1, L0747: 1, L0747: 1, L0748: 1, L0748: 1, L0747: 1, L0748: 1, L0747: 1, L0748: 1, L0747: 1, L0748: 1, L0748: 1, L0747: 1, L0748: 1,	1 and L0752: 1. AR060: 5, AR316: 5, AR089: 4 L0749: 5, L0777: 3, L0526: 2, L0438: 2, H0547: 2, H0295: 1, S0420: 1, S0356: 1, S0354: 1, L0021: 1, H0586: 1, H0497: 1, L0021: 1, H0553: 1, H0169: 1, H0163: 1, S0440: 1, S0422: 1, L0766: 1, L0769: 1, L0764: 1, L0783: 1, L0793: 1, L0657: 1, L0783: 1, L0793: 1, L0657: 1, L044: 1, H0519: 1, S0330: 1, L0439: 1, L0731: 1, S0436: 1, L0592: 1, L0485: 1, L0608: 1, L0601: 1 and S0194: 1.	AR235: 19, AR291: 19, AR289: 17, AR266: 17, AR261: 17, AR238: 17, AR297: 16, AR243: 16, AR296: 15, AR283: 15, AR269: 15, AR287: 14, AR286: 15, AR287: 14, AR275: 14, AR289: 14, AR168: 13, AR182: 13, AR168: 13, AR163: 13, AR161: 13, AR165: 13, AR162: 13, AR165: 13,
Trp-10 to Lys-18, Val-32 to Cys-38, Asp-41 to Thr-47.		
2742	2743	2744
170 - 340	192 - 293	671 - 826
145	146	147
837545	832384	801966
HAQBG57	HAQBY85	HAQBZ15
135	136	137

12, AR271: 12, AR195: 12, AR195: 12, AR255: 12, AR295: 12, AR060: 12, AR039: 11, AR272: 11, AR262: 11, AR293: 11, AR293: 11, AR175:		9, AR196: 8, AR308: 8, AR312: 8, AR312: 8, AR061: 8, AR190: 7, AR200: 7, AR253: 7, AR219: 7, AR219: 7, AR234: 7, AR234:
AR192: AR204: AR260: AR246: AR270: AR193: AR193: AR193: AR166: AR196: AR309: AR309:	AR191: AR173: AR258: AR269: AR205: AR205: AR189: AR189: AR174: AR174: AR174: AR277: AR277: AR277: AR277: AR277: AR277:	AR172: AR168: AR213: AR254: AR313: AR27: AR210: AR180: AR180: AR231: AR231: AR231: AR231:

AR227: 7, AR221: 7, AR211: 7, AR212: 7, AR255: 7, AR104: 7, AR199: 6, AR264: 6, AR199: 6, AR264: 6, AR217: 6, AR214: 6, AR217: 6, AR214: 6, AR216: 6, AR212: 6, AR216: 6, AR222: 6, AR263: 6, AR224: 6, AR263: 6, AR224: 6, AR223: 5, AR230: 5, AR215: 3 H0046: 5, H0295: 2, H0013: 2, H0294: 1, H0619: 1, H0549: 1, H0550: 1, S0222: 1, H0309: 1, H0550: 1, S0222: 1, H0309: 1, H0687: 1, H0688: 1, H0428: 1, H0039: 1, H0424: 1, H0405: 1, H0181: 1, S036: 1, H0163: 1, H0494: 1, S0210: 1, H0593: 1, H0494: 1, S0210: 1, H0593: 1, S0322: 1 and S0042: 1,	H0295: 1, H0318: 1 and L0662: 1.	AR089: 33, AR316: 24, AR060: 16 H0556: 3, S0002: 2, L0518: 2, L0809: 2, H0696: 2, H0265: 1, H0295: 1, S0114: 1, S0134: 1, H0402: 1, H0431: 1, T0039: 1, H0050: 1, H0014: 1, S6028: 1, H0124: 1, H0634: 1, S0440: 1, L0774: 1, L0791: 1, H0660: 1, S0390: 1, S3014: 1, S0032: 1, L0748: 1, L0754: 1, L0750: 1, S0391: 1, H0444: 1, S0026: 1 and L0697: 1.
	2745	2746
	79 - 111	89 - 166
	148	149
	604906	795993
	HAQCE18	НАQСБ94
	138	139

29,	みてててららららまみよみみよみななるでうううううううごうごうごうご
AR316:	9, AR263: 8, AR205: 7, AR204: 6, AR204: 5, AR192: 5, AR192: 5, AR192: 4, AR2312: 4, AR293: 3, AR293: 3, AR293: 3, AR293: 3, AR293: 3, AR293: 3, AR293: 3, AR293: 3, AR293: 3, AR293: 2, AR231: 2, AR231: 2, AR256: 2, AR256: 2, AR256: 2, AR256: 2, AR256: 2, AR285: 2, AR285:
	AR241: 9, AR246: 8, AR246: 8, AR248: 7, 7, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
AF T	######################################
ys-57.	vs-40, he-79, Gly-121, Lys-140, Ser-168.
Pro-52 to Cys-57.	Pro-29 to Lys-40, Glu-66 to Phe-79, Arg-112 to Gly-121, Thr-128 to Lys-140, Lys-158 to Ser-168.
2747 Pr	2748 G G G G G G G G G G G G G G G G G G G
- 518	- 1426
225 -	422 - 1
150	151
560598	
HARAE26	HARAT69
140	141

	103950, 193100, 193400, 200990, 601458, 602096	
	12p13.3-p13.1	
AR284: 2, AR227: 1, AR296: 1, AR232: 1, AR290: 1 S0276: 9, L0439: 4, H0559: 2, H0620: 2, L0507: 2, L0803: 1, S0040: 1, S0001: 1, T0082: 1, H0581: 1, H0078: 1, H0266: 1, S0022: 1, L0375: 1, L0805: 1, L0599:	L0777: 10, H0616: 4, L0758: 4, L0761: 3, L0766: 3, H0648: 3, L0769: 3, L0750: 3, L0755: 3, L0749: 2, L0750: 3, L0755: 3, H0318: 2, H0545: 2, H0373: 2, L0774: 2, L0375: 2, L0577: 2, L0578: 1, H0656: 1, L0682: 1, H0686: 1, L0760: 1, L0762: 1, L0762: 1, L0762: 1, L0769: 1, L0769: 1, L0769: 1, L0769: 1, L0769: 1, L0768: 1, L0666: 1, L0566: 1, L0378: 1, L0666: 1, L0368: 1, L0666: 1, L0779: 1, L0777: 1, L0779: 1, L0777: 1, L0779: 1, L0777: 1, L0779: 1, L0777: 1, L0777	AR060: 5, AR316: 4, AR089: 3 L0747: 25, L0731: 25,
	2749	2750
	435 - 536	1767 - 1916
	152	153
	832380	845848
	HARAZ81	HASAU26
	142	143

				H0013: 23, H0046: 20, H0144:		
				18, L0/69: 13, L0439: 13,		
			-	L0662: 11, L0/58: 10, H0009:		
				9, H0556: 8, T0010: 8, L0764:		
				8, L0748: 8, L0759: 8, S0001:		
				7, L0776: 7, L0659: 7, S0360:		
	-			6, S0007: 6, H0266: 6, L0749:		
				6, L0750: 6, L0594: 6, H0457:		
				5, L0761: 5, L0754: 5, L0589:		
				5, L0592: 5, H0265: 4, H0706:		
				4, S0010: 4, H0264: 4, L0351:		
				4, S0002: 4, L0638: 4, L0806:		
				4, L0805: 4, L0783: 4, L0666:		
•						
				. —		
				3 H0178: 3 L0471: 3 H0012:		
				3 HO620: 3 HO428: 3 HO622:		•
_				2 HOSS1, 2 HOTO, 2 LOTTO.		
			•	3, H0331: 3, H0100: 3, L0770:		
						•••
		-		[3, L0665: 3, H0672: 3, H0696:		
-				[3, L0740: 3, L0608: 3, S0045:		
				2, H0619: 2, H0645: 2, H0351:		
				[2, S0278: 2, S0005: 2, H0599:		
•				2, T0082: 2, H0004: 2, H0052:		
				2, S0250: 2, H0424: 2, H0038:		
				2, H0616: 2, H0623: 2, S0440:		
				2, S0344: 2, L0372: 2, L0771:		
				2, L0768: 2, L0766: 2, L0649:		
				2, L0518: 2, L0809: 2, S0148:		
				2, H0520: 2, S0330: 2, L0602:		
_				2, S0152: 2, H0521: 2, S0406:		
				2, L0612: 2, S0027: 2, S0028:		
				2, L0743: 2, L0744: 2, L0755:		-
				2, S0031: 2, S0434: 2, L0591:		
				2, S0040: 1, H0717: 1, H0716:		
				1, S6024: 1, H0583: 1, H0657:		
				1, H0341: 1, S0282: 1, H0662:		
	_			1, H0638: 1, S0418: 1, S0420:	-	

																																	•				
476:	370:	587:	9622:	090	590:	346:	474:	196:	3530:	3123:	3014:	271:	003:	1. H0213:	3628:	S0364:	366:	135.	0272:	, H0413:	1, H0649:	1, S0208:	L0369:	L0639:	L0374:	L0521:	T0000:	T0636:	L0530:	1, L0663:	1, H0691:	0690	0435:	1, S0328:	, S0037:)756:	0485: 0543:
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	2751	2752	2753	2754	2755	2756	2757	2758
	136 - 252	912 - 669	76 - 168	154 - 243	134 - 271	97 - 213	196 - 312	
	154	155	156	157	158	159	160	161
	654833	834511	654834	598714	847023	588403	598715	780460
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	144	145	146	147	148	149	150	151

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L0794: 3, H0436: 3, S0027: L0779: 3, H0170: 2, H0171: H0341: 2, S0360: 2, H0580: L0717: 2, H0431: 2, H0156 S0010: 2, H0545: 2, S6028: S0003: 2, S0142: 2, S0344: S0422: 2, S0426: 2, L0662: L0766: 2, L0804: 2, L0659: L0666: 2, H0435: 2, S0390:	S0037: 2, S3014: 2, L0741: L0743: 2, L0758: 2, L0759: H0444: 2, L0599: 2, H0624, H0265: 1, H0556: 1, H0657 S0116: 1, S0212: 1, H0663: H0638: 1, S0418: 1, S0356: H0329: 1, S0007: 1, S0045: S0046: 1, H0663: 1, S0045:	1, 2022: 1, S0222: 1, H0581: 1, H0569: 1, H0416: 1, H0622: 1, H0111: 1, H01090:	, H0616: 1, H0551: 1, H0413 , L0564: 1, H0641: 1, H0646 , S0144: 1, H0695: 1, L0598: , L0520: 1, L0630: 1, L0645: , L0521: 1, L0767: 1, L0803: , L058: 1, L0566: 1, L0809: , L0790: 1, L0791: 1, L0792: , L0793: 1, L0665: 1, S0374:	I, L0438: 1, S0126: 1, H0711: 1, H0683: 1, H0648: 1, H0539 1, H0522: 1, S0406: 1, H0631: 1, S0028: 1, L0747: 1, L0731: 1, L0583: 1, S0011: 1, H0136: 1, S0192: 1, S0242: 1, S0276: 1, H0543: 1, H0423: 1, H0422
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	Glu-19 to Pro-27.	
	2759	2760
	158 - 253	237 - 284
	162	163
	847024	847025
	HATCI67	HATCJ27
	152	153

				104311, 109150, 112262, 182600, 182870, 182870, 182870, 245200, 600225, 600225, 601208
	X		16	14q22.1-q24.3
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		Glu-34 to Ser-41.		
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	259 - 381	196 - 333	428 - 568	1066 - 1161
	164	165	166	167
	848564	899395	831097	785628
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	154	155	156	157

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	168	169
	603959	847323
	HATDH23	HATDH55
	158	159

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	75 - 173	23 - 85	232 - 372	139 - 252	154 - 240	230 - 328	2203 - 2328
	170	171	172	173	174	175	176
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	160	161	162	163	164	165	166

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1, S0222: 1, H0409: 1, H0592: 1, H0586: 1, H0486: 1, H0190: 1, H0487: 1, H0156: 1, H0190: 1, T0048: 1, H0156: 1, H0190: 1, H0309: 1, H0183: 1, H0590: 1, H0309: 1, H0183: 1, H0590: 1, H0188: 1, S0338: 1, H0520: 1, H0644: 1, H0188: 1, S0348: 1, H0604: 1, H0644: 1, H0189: 1, S0036: 1, H0644: 1, H0660: 1, H0660: 1, H0660: 1, H0660: 1, H0660: 1, H0660: 1, L0771: 1, L0794: 1, L0640: 1, L0670: 1, L0794: 1, L0640: 1, L0670: 1, L0794: 1, L0640: 1, L0670: 1, L0794: 1, L0640: 1, L0650: 1, L0794: 1, L0660: 1, L0660: 1, H0680: 1, H0435: 1, S0220: 1, H0689: 1, H0435: 1, S0220: 1, H0425: 1, S0032: 1, L0780: 1, S0032: 1, L0780: 1, S0032: 1, H0422: 1, S0032: 1, H0423: 1, S0032: 1, H0422: 1, S0032: 1, H0423: 1, S00	H0265: 1, H0556: 1, S0134: 1, H0580: 1, H0441: 1, T0060: 1 and H0543: 1.	L0777: 6, L0748: 5, L0758: 5, L0439: 4, L0375: 3, L0531: 3, L0663: 3, L0747: 3, L0749: 3, L0596: 3, H0553: 2, H0644: 2, L0809: 2, H0144: 2, L0438: 2, S0330: 2, L0740: 2, L0591: 2, L0591: 2, L0591: 2, L0591: 2, L0591: 2, L0591: 1, H0686: 1, S0376: 1, S0360: 1, H0619: 1, H0549: 1, H0331: 1, T0060: 1, T0114: 1, H0013: 1, S0474: 1, H0421: 1, H0009: 1, H0567: 1, H0046: 1, H0050:
	1, 1,	1
	2775	2776
	140 - 259	228 - 365
	178	179
	877621	542056
	HAWAS41	HAWBA65
	168	169

	· · · · · · · · · · · · · · · · · · ·			4
1, L0471: 1, H0012: 1, H0510: 1, H0188: 1, H0618: 1, H0428: 1, H0059: 1, H0100: 1, S0002: 1, L0769: 1, L0769: 1, L0769: 1, L0669: 1, L0764: 1, L0779: 1, L0662: 1, L0764: 1, L0773: 1, L0657: 1, L0789: 1, L0659: 1, H0520: 1, H0549: 1, H0599: 1, H0549: 1, H0549: 1, L0789: 1, L0789: 1, L0789: 1, L0789: 1, L0789: 1, H0583: 1, H0684: 1, H0555: 1, H0589: 1, L0759: 1, L0780: 1, S0436: 1, L0780: 1, H0543: 1, S0476: 1, S0196: 1, H0543: 1 and H0352: 1.	L0809: 4, L0766: 3, L0439: 3, H0624: 2, H0411: 2, L0794. 2, L0756: 2, L0731: 2, L0005: 1, H0599: 1, L0471: 1, S0051: 1, T0010: 1, H0266: 1, S0150: 1, L0637: 1, L0765: 1, L0803: 1, L0783: 1, H0144: 1, H0672: 1, S0392: 1, L0748: 1, L0779: 1, L0777: 1 and L0759: 1.	L0803: 2, H0411: 1, H0587: 1, L0623: 1, L0652: 1, L0809: 1 and H0710: 1.	H0410: 1 H0410: 2	L0752: 8, L0766: 6, L0754: 5, H0624: 4, H0013: 4, L0775: 4, L0666: 4, H0156: 3, L0803: 3, L0774: 3, L0745: 3, L0740: 3, H0171: 2, H0657: 2, S0003: 2, H0169: 2, L0662: 2, L0805: 2, L0776: 2, L0664: 2, L0665:
			Arg-18 to Arg-24, Arg-40 to Lys-47.	
				2781
	1074 - 1079		101 - 244	646 - 771
	180	181	182	184
	801884	862000	877577	841010
	HBAGH64	HBAGV01	HBAMC50 HBAMC57	
	170	171	172	174

2, H0658: 2, H0670: 2, H0672: 2, H0521: 2, S0404: 2, L0746: 2, L0599: 2, L0608: 2, S0192: 2, H0542: 2, H0686: 1, S0116: 1, H0661: 1, H0662: 1, S0442: 1, S0376: 1, S0444: 1, S0376: 1, S0444: 1, S0376: 1, H0576: 1, H0577: 1, H0374: 1, H0576: 1, H0578: 1, H0578: 1, H0574: 1, H0574: 1, H0576: 1, H0576: 1, H0674: 1, H0622: 1, H0674: 1, H0609: 1, H0404: 1, S0440: 1, H0640: 1, L0796: 1, L0769: 1, L0769: 1, L0769: 1, L0769: 1, L0769: 1, L0773: 1, L0677: 1, L0651: 1, H0660: 1, H0648: 1, S0328: 1, L0663: 1, L0563: 1, L0563: 1, L0563: 1, H0660: 1, H0648: 1, S0328: 1, H0655: 1, L0780: 1, H0655: 1, L0780: 1, H0655: 1, L0780: 1, S0330: 1, H0555: 1, L0780: 1, S0330: 1, H0555: 1, L0780: 1, S0331: 1, S0260: 1, S0436: 1 and H0543: 1.	4: 1	L0766: 11, L0745: 9, L0756: 9, L0752: 5, H0305: 3, S0414: 3, S0422: 3, L0439: 3, H0156: 2, H0374: 2, L0731: 2, L0757: 2, L0757: 2, L0731: 2, L0757: 2, L0756: 1, H0486: 1, H0590: 1, S0222: 1, H0486: 1, H0194: 1, S0388: 1, H0494: 1, S0448: 1, L0655: 1, L0774: 1, L0772: 1, L0768: 1, L0774:
2, H0658: 2 2, H0521: 2 2, L0599: 2 2, H0542: 1, H0661: 1, S0376: 1 1, H0374: 1, H0374: 1, H0123: 1, H0123: 1, H0123: 1, H0622: 1, H0609: 1, L0637: 1, L0637: 1, L0637: 1, L0663: 1, L0663: 1, H0660: 1, H0655: 1, H0655: 1, H0553: 1, H0553: 1, H0553: 1, H0553: 1.	H0374: 1	Pro-44 to Thr-54. L076 3, S04 3, S04 2, H03 2, L07 1, H05 1, S04 1, S04 1, L07
	2782	2783 Pr
	805 - 840	106 - 297
	185	186
	736014	
	HBBBB08	HBBBE83
	175	176

			181031
1, L0375: 1, H0519: 1, S0328: 1, L0779: 1, L0777: 1 and S0106: 1.	H0389: 1	L0803: 4, L0663: 3, S0420: 2, L0666: 2, H0435: 2, L0748: 2, L0777: 2, S0116: 1, S0132: 1, H0370: 1, L0621: 1, S0422: 1, L0664: 1, H0658: 1, S0328: 1, L0749: 1, S0436: 1 and L0097: 1.	AR060: 18, AR316: 11, AR089: 5 L0803: 14, L0744: 13, L0754: 13, H0265: 12, L0731: 11, H0556: 9, H0031: 9, L0740: 9, L0758: 7, S0010: 6, H0038: 6, H0494: 6, L0776: 6, L0747: 5, L0599: 5, S0360: 4, H0253: 4, H0318: 4, H0552: 4, H0574: 4, S0222: 3, H0497: 3, H0574: 3, H0013: 3, S0346: 3, H0574: 3, H0013: 3, S0346: 3, L0663: 3, H0455: 3, L0743: 3, L0663: 3, H0522: 3, L0743: 3, L0652: 3, H0522: 3, L0743: 3, L0751: 3, L0771: 3, H0497: 3, H0574: 3, H0650: 2, H0646: 2, L0608: 3, H0661: 2, S0280: 2, H0570: 2, H0661: 2, S0280: 2, H0574: 2, H0620: 2, H0046: 2, S0250: 2, H0266: 2, H0188: 2, S0250: 2, H0266: 2, H0188: 2, L0456: 2, H0040: 2, L0142: 2, L0456: 2, H0040: 2, L0142: 2, L0456: 2, H0040: 2, L0763: 2, H0100: 2, S0002: 2, L0763: 2,
1,1	H	2,1 2,1 1,1 1,1 1,1 1,1 1,1	HICKSSCHECHCHCHCHCHCHCHCHCHCHCHCHCHCHCHCHCHC
	2784	2785	2786
	199 - 378	96 - 128	124 - 243
	187	888	189
	787308	810227	840585
	HRBMAII	HBCAK10	HBCAK80
	177	178	179

L0772: 2, L0764: 2, L0768: 2, L0804: 2, L0775: 2, L0517: 2, L0783: 2, L0666: 2, L0664: 2, L0665: 2, H0658: 2, H0648: 2, S0330: 2, S3012: 2, S0027: 2,	S0028: 2, L0756: 2, L0596: 2, H0665: 2, H0422: 2, T0002: 1, H0717: 1, H0716: 1, H0294: 1, S0114: 1, S0218: 1, H0656: 1, H0381: 1, H0341: 1, S0212: 1, S0110: 1, S0001: 1, H0483: 1,	H0255: 1, H0669: 1, H0663: 1, H0402: 1, S0418: 1, S0358: 1, S0468: 1, S0140: 1, S0132: 1, H0619: 1, H0645: 1, H0550: 1, H0441: 1, H0431: 1, H0612: 1, H0586: 1, S0005: 1, H0333: 1,	S0414: 1, H0486: 1, H0069: 1, H0156: 1, H0108: 1, H0575: 1, T0082: 1, H0036: 1, H0618: 1, H0581: 1, H0327: 1, H0172: 1, H0010: 1, H0012: 1, H0023: 1,	L0695: 1, L0163: 1, S0051: 1, H0375: 1, H0267: 1, H0271: 1, H0687: 1, H0288: 1, H0292: 1, S0003: 1, H0615: 1, H0428: 1, H0039: 1, H0644: 1, H0424: 1,	L0055: 1, L0455: 1, H0124: 1, H0135: 1, H0163: 1, H0090: 1, H0634: 1, H0087: 1, H0412: 1, H0413: 1, T0004: 1, T0042: 1, L0475: 1, H0509: 1, H0646: 1,	1, H0538: 1, L0638: 1, L0641: 1, L0662: 1, L0551: 1, L0651: 1, L0653:
L077 L080 L078 L066 L066 S033	S002 H066 H071 S011 S01103	H025 H04C S046 H061 H044 H058	S041 H015 T008 H056 H0001	H03: H06: H06: H06: H06: H00:	L00: H01: H06: H04: L04:	50142: L0770: L0646: L0521: L0388: L0774: L0774:

	193300, 193300, 227646, 601154			
	3p25.3-3p24.1			
L0636: 1, L0542: 1, L0782: 1, L0519: 1, L0789: 1, H0144: 1, S0374: 1, S0310: 1, H0519: 1, S0126: 1, H0682: 1, H0672: 1, L0602: 1, H0539: 1, S0380: 1, H0626: 1, S0037: 1, L0748: 1, L0779: 1, L0777: 1, L0752: 1, L0755: 1, S0260: 1, S0434: 1, S0436: 1, S0384: 1 and H0423: 1, S0384: 1 and H0355: 1, L0361: 1, S0260: 1, S0436: 1, H0423: 1, S0384: 1 and H0355: 1, L0361: 1, S0260: 1, H0423: 1, S0384: 1 and H0355: 1, L0361: 1, S0260: 1, H0423: 1, S0384: 1 and H0355: 1, L0361: 1, L0361: 1, S0260: 1, H0423: 1, S0384: 1 and H0355: 1, L0361: 1, L0361: 1, S0260: 1, H0361: 1, S0384: 1 and H0355: 1, L0361: 1, S0384: 1 and H0355: 1, L0361: 1, S0384: 1 and H0355: 1, L0361: 1, L0361: 1, S0384: 1 and H0355: 1, L0361: 1, S0384: 1 and H0355: 1, L0361: 1, S0384: 1 and H0355: 1, S0386: 1, S0384: 1 and H0355: 1, S0384	H0556: 2, H0550: 2, H0581: 2, H0050: 2, H0509: 2, L0593: 2, S0040: 1, H0657: 1, H0656: 1, S0418: 1, H0619: 1, S0278: 1, H0370: 1, H0156: 1, H0052: 1, H0083: 1, H0510: 1, H0033: 1, H054: 1, H0413: 1, T0041: 1, H0132: 1, H0551: 1, H0539: 1, L0438: 1, H0134: 1, S0390: 1, L0439: 1, S0050: 1, H0445: 1, L0601: 1, S0011: 1, H0665: 1 and H0423: 1.	AR089: 39, AR316: 31, AR060: 25 L0439: 10, L0769: 3, H0455: 2, L0526: 2, L0438: 2, L0748: 2, L0751: 2, L0777: 2, L0411: 1, S0110: 1, S0001: 1, L0717: 1, S0222: 1, H0370: 1, L0563: 1, H0041: 1, H0687: 1, H0286: 1, H0413: 1, H0494: 1, S0344: 1, L0770: 1, L0793: 1, H0521: 1, H0187: 1, S0028: 1, L0786: 1 and L0752: 1.	S0334: 1	H0617: 6, L0666: 3, H0188:
	000	Pro-25 to Pro-31.		Pro-20 to Ser-28.
	2787	2788	2789	2790
	320 - 421	281 - 376	225 - 365	298 - 408
	190	191	192	193
	525002	801885	69889	838123
	HBCAQ48	HBCAY17	HBCGE46	HBGBA14
	180	181	182	183

		_																																		
2, H0181: 2, H0196: 1, H0039:	1, L0659: 1, L0809: 1, L0789:	1, L0663: 1, L0603: 1, H0038: 1. S0330: 1, S0378: 1 and	S0406: 1.	AR089: 13, AR316: 12, AR060: 12	H0585: 9, H0617: 8, L0776:	4, L0755: 4, L0758: 4, H0295:	3, S0406: 3, H0484: 2, S0408:	2, H0251: 2, H0546: 2, H0188:	2, H0615: 2, S0440: 2, S0422:	2, L0770: 2, L0764: 2, L0806:	2, H0519: 2, L0748: 2, L0439:	2, L0740: 2, L0747: 2, L0753:	2, L0589: 2, H0265: 1, H0556:	1, H0141: 1, S0134: 1, H0254:	1, H0661: 1, S0356: 1, S0442:	1, S0222: 1, H0370: 1, H0486:	, L0586: 1, S0049: 1, H0562:	1, L0471: 1, H0620: 1, H0024:	1, H0266: 1, H0292: 1, H0213:	1, H0181: 1, H0182: 1, H0606:	1, H0673: 1, S0364: 1, S0366:	1, H0598: 1, H0163: 1, H0087:	1, H0412: 1, H0059: 1, H0100:	1, H0494: 1, H0646: 1, L0763:	1, L0363: 1			1, L0518: 1, L0783: 1, L0809:	1, L0519: 1, L0665: 1, H0702:	1, H0547: 1, H0690: 1, H0684:	1, S0328: 1, S0330: 1, H0539:	S0380: 1, H0696: 1, H0134:	1, H0555: 1, H0447: 1, L0743:	L0750: 1, L0759: 1, H0542:	H0543: 1 and H0423: 1.	L0748: 6, L0740: 4, L0731:
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				194 161 - 328 2791				-																												195 109 - 132 2792
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	102200, 106100, 131100, 131100, 131100, 133780, 147050, 153700, 161015, 164009, 168461, 188461, 180840, 191181, 193235, 209901, 232600, 259770, 600045, 600319, 600528,		
	11913	16	=
4, L0717: 3, L0776: 3, L0439: 3, L0747: 2, L0749: 3, H0617: 2, L0774: 2, L0663: 2, L0779: 2, L0758: 2, H0402: 1, S0360: 1, H0441: 1, H0156: 1, S0024: 1, H0428: 1, H0181: 1, L0455: 1, L0772: 1, L0771: 1, L0766: 1, L0655: 1, L0661: 1, L0789: 1, L0532: 1, H0593: 1, L0754: 1, L0754: 1, L0758: 1, L07	AR089: 13, AR316: 11, AR060: 10 H0615: 2 and H0606: 1.	AR089: 32, AR316: 27, AR060: 23 H0617: 1	AR089: 43, AR316: 35, AR060: 28 H0617: 25, H0422: 5, H0181: 4, L0743: 4, H0549: 3, L0666: 3, H0658: 3, H0436: 3, H0543: 3, H0583: 2, H0402: 2, S0354: 2, H0486: 2, H0688: 2, S0366: 1, S0358: 1, S0360: 1, S0366: 1, S0358: 1, S0360: 1, H06643: 1, H0550: 1, H0618: 1, H0485: 1, H0590: 1, H0618: 1, H04857:
3 3 7 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Ser-21 to Ser-26, Glu-32 to Pro-37.	Glu-28 to Phe-33, Ala-56 to Gln-62.	Gly-27 to Met-38.
	2793	2794	2795
	86 - 199	90 - 431	163 - 279
	196	197	861
	793803	839192	840586
	НВGFQ34	HBGML95	HBGMT60
	186	187	88

	AR213: 37, AR241: 31, AR089: 18, AR198: 17, AR204: 16, AR271: 15, AR229: 17, AR186: 16, AR229: 15, AR185: 15, AR229: 13, AR240: 13, AR300: 13, AR240: 13, AR300: 13, AR240: 13, AR300: 13, AR275: 13, AR293: 12, AR316: 12, AR293: 12, AR316: 12, AR293: 11, AR194: 11, AR293: 11, AR194: 11, AR271: 11, AR194: 11, AR271: 11, AR258: 11, AR260: 10, AR274: 10, AR272: 11, AR258: 11, AR260: 9, AR270: 9, AR246: 9, AR270: 9, AR260: 9, AR291: 8, AR292: 8, AR291: 7, AR292: 8, AR294: 7, AR295: 7, AR294: 7, AR296: 7, AR294: 7, AR296: 7, AR296: 7, AR296: 7, AR296: 6,
1,1 1,1 1,1 1,1 1,1 1,1 1,1 1,1 1,1	Lys-44 to Thr-50, AR Gln-111 to Gly-118, AR
	53 - 541 2796
	HBHAA53 603183 199
	H 681

AR315: 6, AR227: 6, AR179: 6, AR266: 6, AR244: 6, AR248: 5, AR298: 5, AR295: 5, AR183: 5, AR289: 5, AR259: 4, AR232: 4, AR284: 4, AR283: 4, AR314: 4, AR253: 3, AR281: 2, AR265: 2	AR089: 12, AR316: 10, AR060: 8 L0751: 8, L0749: 3, H0179: 2, L0766: 2, L0789: 2, L0740: 2, L0747: 2, L0756: 2, L0731: 2, T0002: 1, H0333: 1, S0049: 1, H0024: 1, H0316: 1, L0369: 1, L0771: 1, L0806: 1, L0776: 1, L066: 1, L0665: 1, H0521: 1, L0779: 1 and H0343: 1.	
4444444		Tyr-15 to Asn-21. Pro-59 to Ala-64.
	2797	2798
	202 - 258	17 - 109
	500	202
	840354	596805 897484
	HBIAU43	HBIBB20
	190	191

·	102700, 102700, 602025
	20q11.21- q13.11
1, H0622: 1, L0194: 1, H0135: 1, H0412: 1, H0623: 1, H0059: 1, L0351: 1, T0042: 1, H0561: 1, S0294: 1, L0640: 1, L4747: 1, L5775: 1, L5565: 1, L0800: 1, L0744: 1, L0748: 1, L0768: 1, L0774: 1, L0776: 1, L0559: 1, L0559: 1, L0559: 1, L0559: 1, L0559: 1, L0579: 1, L0559: 1, L0566: 1, L0664: 1, H0520: 1, H0547: 1, S0378: 1, S0190: 1, S0406: 1, H0436: 1, L0748: 1, L0748: 1, L0759: 1, L0601: 1, L0366: 1 and H0423: 1.	AR186: 30, AR061: 21, AR298: 20, AR314: 17, AR055: 17, AR104: 16, AR229: 15, AR259: 15, AR249: 15, AR284: 14, AR249: 15, AR284: 14, AR257: 14, AR292: 13, AR052: 12, AR233: 12, AR052: 11, AR182: 11, AR185: 11, AR280: 10, AR241: 10, AR280: 10, AR248: 9, AR289: 10, AR248: 9, AR293: 9, AR248: 9, AR293: 9, AR248: 9, AR293: 9, AR275: 9, AR293: 7, AR267: 7, AR238: 7, AR267: 7, AR238: 7, AR266: 6, AR312: 6, AR213: 6, AR312: 6, AR313: 6,
	Pro-67 to Ser-78, Arg-114 to Ser-125, Ser-147 to Ala-155. A A A A A A A A A A A A A A A A A A A
	2800
	655 - 1206
	845743 203
	HBIBF26
	193

														•																					
	389: 0, AR206: 0, 353: 6, AR269: 6,	6, AR198:	5, AR290:	5, AR316:	AR219: 5, AR000: 5, AR296: 5, AR310: 5,	5, AR281:	5, AR274:		AR309: 4, AR204: 4,			AR218: 4, AR263: 4,	AR256: 4, AR177: 4,	AR202: 4, AR246: 4,	AR168: 3, AR205: 3,	AR165: 3, AR166: 3,	AR164: 3, AR261: 3,		AR197: 2, AR176: 2,	. •		• •		AR196: 1, AR210: 1	L0659: 6, H0657: 4, H0521:	4, L0439; 4, L0/45; 4, L0/59; 4, 110556; 2, 50250; 3, 1,0751;	4, H0330; 3, 30300; 3, E0701;] 2 1 0662; 3 1 0766; 3 1 0809;]	3, H0549; 2, H0392; 2, H0253;	2, H0581: 2, H0620: 2, H0051:	2, H0551: 2, H0494: 2, L0770:	2, L0794: 2, L0649: 2, L0665:	2, H0520: 2, S0032: 2, L0741:	2, L0743: 2, L0748: 2, L0747:	L0779: 2, L0758: 2, L0605:	2, H0650: 1, H0484: 1, H0254: 1, H0402: 1, S0358: 1, H0580:
AR243:	AR089: AR053:	AR270:	AR2	AR2	ARZ	ARZ	ARC	ARC	AR	AR	ARC	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR		4, I	4, 1	7,7	2,1	2,1	[2, I	2,1	2,1	2,1	2,1
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	Ser-11 to Gln-18.	Gly-23 to Thr-33.	
	2801	2802	2803
	2 - 76	71 - 187	112 - 219
	204	205	206
	852045	598719	580807
	HBIBM33	HBIBN67	нвів (69
	194	195	196

											112262, 182600,	182870, 182870,	182870, 600225,	000223	601011, 601011,	601011, 601011,	001/08, 001843																
		<u>e</u>									14q22.1			-	19p13			•••	- 2		-:-		<u></u>				•••						
H0265: 1, S0049: 1 and S0428: 1.	S0049: 1	AR089: 4, AR316: 4, AR060: 3	L0741: 6, L0439: 5, L0742: 4, L0352: 3, S0010: 2, L0764:	2, L0768: 2, L0717: 1, H0013:	1, H0318: 1, S0049: 1, H0052: 1 H0009: 1, S0036: 1, H0135:	1, H0087: 1, L0351: 1, L0749:	1, L0755: 1, S0031: 1 and	L0594: 1.	S0049: 1	S0049: 1, H0416: 1 and H0316: 1.	S0049: 1				S0358: 10, L0439: 5, H0009:	4, S0360: 3, S0408: 3, H0585:	2, H0657: 2, S0420: 2, S0442:	2, H0734: 2, H0318: 2, H0052	2, H0373: 2, H0510: 2, H0266:	2, H0708: 2, H0445: 2, H0656:	1, H0341: 1, H0484: 1, H0662	1, H0730: 1, H0722: 1, H0728:	1, H0392: 1, H0587: 1, H0559:	1, H0486: 1, S0010: 1, H0581:	1, S0049: 1, H0251: 1, H0545:	1, H0687: 1, H0644: 1, H0617:	1, S0364: 1, H0494: 1, H0633:	1, H0647: 1, S0144: 1, S0422:	1, H0529: 1, L0761: 1, L0773:	1, H0693: 1, H0593: 1, S0126:	, H0682:	1, H0732: 1, L0742: 1, L0743:	1, L0/45: 1, S0436: 1, L0594:
		7	7																														
	2804	2805						•	2806	2807	2808				2809																		
	27 - 74	235 - 243							234 - 305	65 - 166	146 - 202				190 - 252														-				
	207	208							209	210	211				212																		
	612783	824258						•	590280	608320	688861				637516																		
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	197	861							100	200	201	1			202																		

	141/50, 141800, 141800, 141800, 141800, 141850, 141850, 141850, 141850, 156850, 186580, 191092, 600173, 601785							164500, 176880, 232500, 600151, 600795
	16p13.3							3p12-p11.1
1, S0011: 1, H0542: 1 and H0352: 1.	AR060: 6, AR316: 4, AR089: 2 L0438: 6, L0741: 5, H0556: 4, S0051: 4, S0036: 4, L0774: 4, H0622: 3, L0769: 3, H0619: 2, H0261: 2, H0318: 2, H0194: 2, L0471: 2, L0757: 2, L0758: 2, L0593: 2, H0624: 1, H0265: 1, H0650: 1, H0657: 1, S0222: 1, H0455: 1, H0559: 1, H0055: 1, H0455: 1, H0559: 1, H0056: 1, H0455: 1, H0594: 1, H0644: 1, H0083: 1, H0594: 1, H0615: 1, H0083: 1, H0594: 1, H0615: 1, H0494: 1, S0450: 1, S0144: 1, L0770: 1, L4777: 1, L0639: 1, L0761: 1, L0775: 1, L0839: 1, L0635: 1, L0788: 1, S0428: 1, L0748: 1, S0011: 1 and H0136: 1.	H0318: 1 and S0027: 1.	H0583: 1, H0318: 1 and H0309: 1.	AR089: 5, AR316: 5, AR060: 5	L0766: 5, H0318: 1 and L0649: 1.		H0318: 1 and S0037: 1.	AR089: 17, AR316: 13, AR060: 9 L0803: 12, L0766: 10,
			Ser-31 to Lys-39.			Asp-16 to Glu-27.	Lys-25 to Trp-32.	
	2810	2811	2812	2813		2814	2815	2816
	329 - 370	274 - 288	90 - 209	173 - 199		50 - 172	146 - 334	55 - 165
	213	214	215	216		217	218	219
	841235	603529	626591	209095		581104	609859	847332
	HBJAC40	HBJAV56	HBJAY14	нвлв069		HBJBR40	HBJCH46	HBJCR17
	203	204	205	506		207	208	209

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H0457: 8, L0777: 8, L0759: 8, L0752: 7, L0754: 6, H0521: 5,	L0439: 4, L0740: 4, H0170: 3, S0222: 3, H0587: 3, H0090: 3,	L0770: 3, L0805: 3, L0659: 3,	H0686: 2, L0415: 2, S0116: 2, H0407: 2, L0005: 2, H0486: 2.	H0013: 2, H0036: 2, H0318: 2,	H0581: 2, H0251: 2, H0373: 2,	H0032: 2, H0038: 2, H0040: 2, H0634: 2, H0551: 2, L0520: 2,	L0375: 2, L0809: 2, L0664: 2,	L0665: 2, L0438: 2, H0547: 2,	H0555: 2, L0779: 2, L0755: 2,	L0731: 2, L0758: 2, L0596: 2,	LOS92: 2, LO004: 2, LOS94: 2, CO040: 1 SO134: 1 H0650: 1.	H0657: 1, 10811: 1, H0483: 1,	H0638: 1, S0354: 1, S0358: 1,	S0360: 1, H0329: 1, S0045: 1,	S0046: 1, L0717: 1, H0351: 1,	H0497: 1, H0574: 1, H0632: 1	1, L0021: 1	 H0569: 1, H0123: 1, 50031: 1, H0260: 1 S0003: 1 S0214: 1		H0622: 1, H0031: 1, H0553: 1	H0124: 1, H0068: 1, H0135: 1	H0591: 1, H0616: 1, H0488: 1	H048/: 1, H0412: 1, S0440: 1, H0130: 1 H0652: 1. L0598: 1,	L0369: 1, L0638: 1, L0646: 1,	L0521: 1, L0662: 1, L0388: 1,	L0804: 1, L0775: 1, L0655: 1,	L0517: 1, L0791: 1, L0793: 1,	L0666: 1, L0663: 1, H0520: 1,	H0519: 1, S0126: 1, H0690: 1,	H0659: 1, H0660: 1, H0696: 1 H0710: 1, S0152: 1, H0696: 1
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S0044: 1, S0404: 1, S0406: 1, H0436: 1, S0028: 1, L0743: 1, L0747: 1, L0750: 1, L0786: 1, H0445: 1, H0136: 1, S0196: 1	AR223: 18, AR221: 16, AR224: 15, AR217: 14, AR168: 12, AR217: 10, AR215: 9, AR216: 9, AR222: 9, AR216: 9, AR221: 10, AR214: 8, AR216: 9, AR214: 8, AR311: 8, AR312: 8, AR213: 7, AR191: 7, AR191: 7, AR192: 6, AR263: 6, AR293: 6, AR293: 5, AR196: 5, AR293: 5, AR196: 5, AR293: 5, AR293: 5, AR293: 5, AR293: 5, AR293: 4, AR296: 4, AR299: 4,	4, AR181:
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AR257. 4 AR218. 4.				3, AR060:	AR226: 3, AR203: 3,		3, AR195:	3, AR200:	2, AR205:	AR104: 2, AR261: 2,	2, AR232:	2, AR267:	2. AR233:	2, AR211:	2, AR183:	1, AR210:	AR247: 1	L0766: 9, L0731: 9, L0754:	7, L0777: 7, L0439: 6, H0551:	5, L0758: 5, L0759: 5, L0666:	4, H0521: 4, L0747: 4, L0592:	4, L0608: 4, H0423: 4, H0657:	3, H0662: 3, H0013: 3, H0038:	[3, H0494: 3, L0438: 3, L0756:	[3, L0779: 3, L0752: 3, L0755:]	3, S0358: 2, H0486: 2, H0251:	2, S0003: 2, H0090: 2, L0598:	[2, L0764: 2, L0794: 2, L0776:]	2, L0663: 2, L0665: 2, S0374:	2, H0547: 2, H0682: 2, H0648:	2, S0152: 2, H0555: 2, H0478:	2, L0740: 2, L0591: 2, H0543:	2, S0412: 2, S0114: 1, L0760:
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S0212: 1, S0418: 1 S0360: 1, T0008: 1 H0580: 1, S026: 1 S6026: 1, S0222: 1 H0069: 1, L0021: 1 H0150: 1, L0471: 1 H0271: 1, S0312: 1 H0688: 1, H0266: 1 H0688: 1, H0428: 1 H0646: 1, S0426: 1 L0769: 1, L0637: 1 L0767: 1, L0659: 1 H0666: 1, S0426: 1 H0144: 1, H0701: H0593: 1, H0658: 1 H0666: 1, S0328: S0378: 1, H0658: 1 H0666: 1, S0328: S0378: 1, H0696: S0028: 1, L0589: 1 H0445: 1, L0589: H0445: 1, H0696: S0028: 1, L0589: 1 H0445: 1, L0589: H0445: 1, L0589: S0028: 1, L0589: 1	1 and H0506: 1. H0009: 69, H0144: 63, S0007: 60, H0265: 59, H0556: 56, H0013: 55, H0178: 55, L0748: 55, T0010: 48, H0052: 40, S0010: 38, H0100: 38, L0595: 35, H0124: 32, S0027: 30, H0266: 29, H0341: 28, H0521: 27, L0593: 27, H0031: 26, H0040: 23, L0588: 23, L0591: 23, T0049: 22, H0351: 22, H0083: 22, S0036: 22, H0090: 22, H0318: 20, S0038: 18, H0024: 17, L0439: 17, H0295: 16, S0046: 16, S0049:
	1
	3 2818
	221 323 - 403
	HBJCW24 805962
	211

16, S0011: 16, S0045: 15, H0494: 15, S0031: 15, L0594: 15, H0543: 15, H0171: 14, H0123: 14, L0747: 14, L0608: 14, S0040: 13, H0046: 13,	L0754: 13, H0584: 12, H0051: 12, S0022: 12, H0135: 12, L0740: 12, L0596: 12, L0601: 12, H0170: 11, H0294: 11, S0278: 11, H0327: 11, H0612: 11, H0551: 11, S0002: 11, S0126: 11, T0039: 10, S0144: S0126: 11, T0039: 10, S0144: S0126: 11, S0002: 1	10, H0333: 9, H0156: 9, N0006: 9, H0050: 9, S6028: 9, T0006: 9, H0617: 9, H0087: 9, H0522: 9, S0037: 9, S0212: 8, H0125: 8, H0251: 8, S0051: 8, H0039: 8, H0063: 8, H0264: 8,	NOZIU: 8, SUZOG: 8, LO/45: 6, H0352: 8, T0002: 7, H0492: 7, S0346: 7, H0213: 7, H0547: 7, H0519: 7, L0744: 7, L0605: 7, L0362: 7, S0222: 6, H0041: 6, H0172: 6, H0081: 6, H0188: 6, H0286: 6, T0041: 6, T0042: 6, H0560: 6, H0539: 6, L0602: 6,	S0032: 6, H0542: 6, H0008: 6, S0132: 5, H0486: 5, H0014: 5, H0594: 5, H0290: 5, S0214: 5, H0520: 5, H068: 5, H0591: 5, H0520: 5, L0599: 5, S0026: 5, H0136: 5, S0276: 5, H0167: 4, S0116: 4, S0282: 4, S0356: 4, H0369: 4, H0261: 4, S0356: 5, S0356:	H00101: 4, H0427: 4, H0575: 4, H0018: 4, H0596: 4, H0544: 4, H0546: 4, T0003: 4, H0284: 4, H0428: 4, H0413: 4, H0056: 4, S0344: 4, S3014: 4, L0750: 4, L0757: 4, S0260: 4, L0587: 4,
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L0590: 4, L0603: 4, H0140: 3, S6024: 3, S0001: 3, H0484: 3, H0228: 3, H0393: 3, S0300: 3, H0431: 3, H0587: 3, T0040: 3, H0250: 3, T0048: 3, H0209: 3, H0204: 3, H0209: 3, H0204: 3, H0209: 3, H0244: 3, H0208: 3, H0208: 3, H0208: 3, L0755: 3, H0445: 3, L0597: 3, L0597: 3, S0400: 2, H0255: 2, H0423: 3, S0354: 2, S0358: 2,	S0360: 2, H0438: 2, H0069: 2, S0280: 2, H0036: 2, H0004: 2, H00253: 2, S0182: 2, H0581: 2, H0421: 2, H0545: 2, H0150: 2, H0201: 2, H0267: 2, H0179: 2, H0292: 2, S0003: 2, H0252: 2, H0630: 2, H0166: 2, S0364: 2, H0639: 2, H0616: 2, H0616: 2, H0412: 2, H0059: 2, H0661: 2, H0435: 2, H0060: 2, L0435: 2, H0060: 2, H0640: 2, H0640: 2, H0660: 2, H0640: 2, H0060: 2, L0435: 2, H0060: 2, L0426: 2, L0426	H0517: 2, 10202: 2, 20053: 2, 20152: 2, 20064: 2, 20053: 2, 20152: 2, 20064: 2, 20152: 2, 20152: 2, 20060: 2, 20134: 1, 20156: 2, 20600: 2, 20134: 1, 20168: 1, 20160: 1, 20029: 1, 20468: 1, 10459: 1, 10459: 1, 20468: 1, 10229: 1, 10151: 1, 20468: 1, 20229: 1, 10259: 1, 20268: 1, 20269: 1, 20268: 1, 20269:
1.0590: 86024: H0228: H0431: H0309: T0115: 80388: H0102: R0423: H0423:	80360: 80280: 80280: H0253: H0011 H0375 H0292 H0698 H0616 H0616	H0517 S01527 L0731 L0731 H0216 H0483 H0483 S0408 S0408 S0468 H0208 H0497 H0497 H0497

			106165, 117700, 117700, 150210, 169600, 180380, 180380, 180380,
·	22		3q21
H0042: 1, H0505: 1, H0085: 1, H00457: 1, H0597: 1, H0457: 1, H0199: 1, H0199: 1, H0015: 1, H0095: 1, H0095: 1, H0095: 1, H0097: 1, H0095: 1, H0007: 1, H0067: 1, H0059: 1, H0059: 1, H0067: 1, H0059: 1, H0059: 1, H0067: 1, H0059: 1, H0059: 1, H0059: 1, H0064: 1, H0533: 1, H0111: 1, H0081: 1, H0032: 1, H0011: 1, H0018: 1, H0036: 1, H0059: 1, H0079: 1, S0386: 1, S0112: 1, S0016: 1, H0036: 1, S0352: 1, S0370: 1, S0306: 1, H0038: 1, S0372: 1, S0372: 1, S0374: 1, S0374	L0731: 2, H0635: 1, H0318: 1, H0581: 1, H0582: 1, L0766: 1, L0783: 1, L0809: 1, L0666: 1, L0665: 1, S0052: 1, L0439: 1, L0740: 1, L0747: 1, L0780: 1, L0758: 1, L0758: 1, L0366: 1 and S0424: 1.	H0318: 1 and L0779: 1. H0318: 1	80380: 5, H0624: 4, H0547: 4, H0305: 3, L0740: 3, L0754: 3, L0779: 3, H0171: 2, H0657: 2, S0376: 2, S0360: 2, S0222:
			Ser-18 to Asn-23, Leu-34 to Ile-39.
	2819	2820	2822
	173 - 202	265 - 318 141 - 218	56 - 112 48 - 185
	222	223 224	225
	603516	604907 600395	737793 866158
·	HBJDC57	1 - 1 - 1	HBJEE51 HBJEL21
·	212	213	215

190000, 203500, 232050, 276902, 600882, 601199, 601199, 601471, 601682	
2, H0625: 2, L0766: 2, L0659: 2, L0636: 2, H0520: 2, L0748: 2, H0445: 2, L0604: 2, H0653: 2, L0600: 2, S0114: 1, S0134: 1, L0808: 1, H0661: 1, S0442: 1, H00438: 1, H00498: 1, H00418: 1, H0058: 1, H0052: 1, S0028: 1, H0388: 1, H0067: 1, H0057: 1, H0551: 1, H0056: 1, L0749: 1, L0364: 1, L0809: 1, L0655: 1, L0749: 1, L0777: 1, L0777: 1, L0757: 1, L0594: 1, L0777: 1, L0757: 1, L0594: 1, L0362: 1, S0028: 1, L0594: 1, L0594: 1, L0362: 1, S0028: 1, L0777: 1, L0594: 1, L0362: 1, L0577: 1, L0594: 1, L0362: 1, S0028: 1, L0594: 1, L0362: 1, S0028: 1, L0594: 1, L0362: 1, S0028: 1, L0362: 1,	AR089: 10, AR316: 9, AR060: 8
	-
	2824
	1663 - 1686
	227
	836997
	HBJFH84
	217

	109690, 109690, 131400, 138491, 138491, 138491, 154500, 180071, 181460, 222600, 272750, 600807, 601596, 602089
	5432
1, S0448: 1, S0440: 1, S0142: 1, H0529: 1, L0763: 1, L0804: 1, L0775: 1, L0651: 1, L0806: 1, L0805: 1, L0790: 1, L0791: 1, L0664: 1, H0701: 1, H0593: 1, H0670: 1, H0539: 1, S0406: 1, S04	H0013: 23, L0747: 20, H0144: 17, L0731: 17, L0439: 12, L0662: 11, L0769: 10, S0360: 6, S0007: 6, L0776: 6, L0594: 6, H0009: 5, L0758: 6, L0594: 6, H0009: 5, L0758: 4, L0764: 4, L0806: 4, L0805: 4, L0764: 4, L0806: 4, L0805: 3, H0244: 3, L0471: 3, H0428: 3, H0622: 3, H0696: 3, L0750: 3, H0622: 3, H0696: 3, L0750: 3, L0608: 3, S0046: 2, H0645: 2, H06351: 2, S0005: 2, L0700: 2, L0649: 2, L071: 2, L0768: 2, L0649: 2, L0809: 2, S0148: 2, H0672: 2, S0330: 2, S0040: 1, S6024: 1, H0294: 1, S0282: 1, H0662: 1, H0638: 1, S0418: 1, S6024: 1, H0294: 1, S0220: 1, H0649: 1, S0045: 1, H0649: 1, H0294: 1, S0220: 1, H0662: 1, H0638: 1, S0418: 1, S0278: 1, S0614: 1, S0220: 1, H0441: 1, H0455: 1, H0588: 1, L0698: 1, S0360: 1, H0388: 1, S0278: 1, S0614: 1, S0220: 1, H04486: 1, T0060: 1, H0156: 1,
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	8 2825
	228 264 - 308
	14 845228
	218 HBJFJ14

AR309: 21, AR271: 17, AR282: 16, AR272: 16, AR197: 15, AR060: 15, AR245: 14, AR162: 14, AR161: 14, AR291: 13, AR291: 13, AR198: 13, AR163: 13, AR198: 13, AR163: 12, AR165: 12, AR166: 12, AR289: 12, AR166: 12, AR299: 12, AR299: 11, AR209: 11, AR209: 11, AR209: 11, AR209: 11, AR201: 11,
Gly-15 to Asp-25.
180 - 359 2826
1306969 229
219 HBJF126

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	-														_																			
4: 10, 3: 10,																																		16: 7, 36: 7,
10, AR214: 10. AR053:	10, AR192: 10, AR189:	10, AR188: 10, AR188: 10, AR221:	10, AR274:	10, AR250:	10, AR264:	9, AK232. 0 AR238:	9, AR237:	9, AR196:	9, AR180:	9, AR199	9, AR253	9, AR257:	9, AR225	9, AR270	8, AR267:	8, AR243:	8, AR20	8, AR10	8, AR231:	8, AR224:	8, AR290	8, AR210:	8, AR173:	8, AR235;	8, AR033:	8, AR226:	8, AR200:	7, AR061:	7, AR286:	7, AR179:	7, AR233:	7, AR218:	7, AR170:	7, AR216: 7, AR256:
AR263: 1 AR177: 1						AK288: AD220:				AR247:	AR191:	AR185:	AR242:	AR297:	AR190:	AR285:					AR255:	AR236:	AR239:	AR219:	AR287:	AR283:	AR277:	AR211:	AR172:	AR039:	AR168:	AR300:	AR262:	AR232: AR228:
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AR311: AR299: AR313: AR203: AR171: AR227:	AR256: AR222: AR215: L0731: 7 5, L07766	3, 50026 2, 20474 2, L0768 2, L0653 2, H0701 2, L0755	1, S0444 1, H0549 1, H0331 1, H0706 1, H0123	1, H051C 1, H0553 1, S0366 1, H056C 1, L077C	1, L0776 1, L07661 1, L0783 1, S0052 1, H0677 1, L0748 1, L0758 1, H059
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	9455.1-455.3	9q33.1-q33.3				11q22				a.							
	L0777: 4, H0318: 2, L0439: 2, L0758: 2, H0170: 1, L0768: 1, L0549: 1, L0665: 1, L0438: 1, L0611: 1, L0740: 1, L0756: 1 and L0593: 1.	L0777: 4, H0318: 2, L0439: 2, L0758: 2, H0170: 1, L0768: 1, L0549: 1, L0665: 1, L0438: 1, L0611: 1, L0740: 1, L0747: 1, L0756: 1 and L0593: 1.	H0369: 1 and H0318: 1.	L0748: 4, L0596: 2, S0442: 1, H0318: 1, H0178: 1, H0674: 1, S0438: 1, H0647: 1, L0800: 1, L0521: 1, L0766: 1, L0528:	1, H0684: 1, S0406: 1, H0555: 1 and L0752: 1.	AR060: 17, AR316: 13,	AR089: 8 L0803: 2, L0666: 2, H0539: 2, H0402: 1, H0450: 1, H0370:	1, 20162: 1, 110210: 1, 20201: 1, L0804: 1, L0521: 1, L0766: 1, L0804:	1, L0651: 1, L0526: 1, L0789:	1, L0756: 1, L0779: 1, L0777:	1, L0731: 1, L0758: 1, L0759: 1 and L0592: 1.	AR089: 28, AR316: 23,	AR060: 17 H0318: 1	H0318: 1, H0150: 1 and L0385: 1.	11	AR060: 4, AR316: 4,	L0667: 2, S0114: 1, H0351: 1, H0318: 1, H0615: 1 and
Gly-15 to Asp-25.						Lys-65 to Thr-71.						Arg-8 to Asn-13.					
5195	2827	2828	2829	2830		2831						2832		2833	2834	2835	
851 - 1030	246 - 350	246 - 350	423 - 539	354 - 449		14 - 247						181 - 222		189 - 392	121 - 177	200 - 265	
2598	230	231	232	233		234	1					235		236	237	238	
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L0764: 1.	AR060: 4, AR316: 4, AR089: 4	L0667: 2, S0114: 1, H0351:	1, H0318: 1, H0615: 1 and 1, H0764: 1.	H0318: 1 and H0038: 1.	AR316: 9, AR089: 9,	AR060: 9	H0556: 12, L0439: 12,	H0266: 11, L0755: 9, H0264:	8, L0766: 8, H0521: 8, L0731:	8, H0634: 7, S0028: 7, L0752:	7, H0012: 6, L0748: 6, L0666:	5, L0754: 5, L0758: 5, S0360:	4, H0623: 4, L0665: 4, L0740:	4, L0750: 4, L0756: 4, L0591:	4 H0265: 3 H0638: 3 H0620:	3, H0135; 3, H0163; 3, H0040;	3 H0551: 3, H0100: 3, S0002:	3, L0769: 3, S0052: 3, H0144:	3, S0126: 3, S3014: 3, H0650:	2, H0305: 2, S0278: 2, H0333:	2, H0632: 2, H0485: 2, T0040:	2, H0013: 2, H0069: 2, H0635:	2, H0575: 2, H0318: 2, H0009:	2, H0123: 2, S0022: 2, H0622:	2, H0553: 2, H0617: 2, H0068:	2, H0038: 2, H0616: 2, H0268:	2, H0561: 2, S0370: 2, H0130:	2, S0142: 2, L0770: 2, L0771:	2, L0662: 2, L0657: 2, L0664:	2, H0518: 2, S0152: 2, H0522:	2, H0555: 2, H0576: 2, L0749:	2, L0605: 2, L0599: 2, H0624:	1, H0140: 1, H0685: 1, H0717	1, H0295: 1, H0662: 1, H0459	1, S0418: 1, S0420: 1, S0358:	1, S0376: 1, S0408: 1, S0476: 1 H0545: 1 H0550:
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		Arg-17 to Glu-31.	4	Arg-23 to Gly-29, A Ser-52 to Glu-74, A
	2839	2840	2841	2842
	236 - 325	228 - 338	284 - 340	309 - 1355
	242	243	244	245
	743181	612069	839735	837242
	HBJJA26	HBJJX02	нвл.н78	HBJND04
	232	233	234	235

H0046: 16, L0803: 14, L0439: 5, L0794: 4, L0804: 4, L0809: 3, L0777: 3, L0759: 3, S0358: 2, H0318: 2, H0553: 2, H0169: 2, L0770: 2, L0771: 2, L0750: 2, L0752: 2, L0731: 2, S0436: 2, S0134: 1, H0656: 1, S0436: 1, S0444: 1, S0046: 1, H0393: 1, S0300: 1, S0222: 1, H0441: 1, H0575: 1, T0082: 1, L0471: 1, H0039: 1, H0528: 1, H0032: 1, H0674: 1, H0598: 1, H0268: 1, L0774: 1, L0548: 1, L0791: 1, L0654: 1, L0548: 1, L0782: 1, L0519: 1, L0548: 1, L0782: 1, L0639: 1, H0593: 1, L0782: 1, L0639: 1, L0548: 1, L0782: 1, L0519: 1, L0548: 1, L0782: 1, L0639: 1, H0593: 1, L0782: 1, L0639: 1, H0593: 1, L0782: 1, L0639: 1, H0593: 1, L0788: 1, H0547: 1, H0593: 1, L0788: 1, L068: 1, H0553: 1 and S0242: 1.	H0318: 1, H0264: 1 and S0426: 1.	S0366: 1, L0769: 1 and L0790: 1.	AR089: 13, AR316: 11, AR060: 8 H0733: 6, S0364: 6, S0366: 6, H0729: 4, L0623: 3, H0708: 3, L0777: 3, H0728: 2, H0706: 2, H0644: 2, L0794: 2, L0750: 2, S0116: 1, S0212: 1, H0586: 1, L0622: 1, H0122: 1, H0581: 1, S0049: 1, H0018: 1, L0644: 1, L079: 1,
Ser-76 to Lys-82, Glu-88 to Pro-94, Pro-111 to Gly-127, Glu-130 to Gly-136, Ile-140 to Gly-136, Irp-176 to Asn-183, Lys-201 to Gly-206, Leu-225 to Gly-230, Sultantial to Gly-230, Leu-237 to Leu-251, Val-267 to Asp-274.	Glu-29 to Asn-34.		His-41 to Gly-46.
	2843	2844	2845
	20 - 124	222 - 311	56 - 232
	246	247	248
	612785	827276	827277
	HBJND57	HBKDF66	HBKEA94
	236	237	238

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S0366: 1 and H0598: 1.	AR196: 7. AR235: 6.	6, AR309:	5, AR311:	AR313: 5, AR312: 5,	 5, AR189:	5, AR183:	5. AR219:	4. AR162:	4, AR263:	4, AR253:	4, AR163:	4, AR178:	4, AR214:		4, AR224:	4, AR166:		AR275: 4, AR270: 4,			3, AR262:	3, AR177:	3, AR288:		AR060: 3, AR182: 3,	AR258: 3, AR245: 3,	AR201: 3, AR316: 3,	AR293: 3, AR218: 3,	3, AR195:	3, AR294:	3, AR274:	AR179: 3, AR181: 3,
	Thr. 115 to Ala-123																															
2846	2847	1407																														
47 - 295	010	441 - 017																														
240	250	007																														
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020	607	240						-																 _								

AR277: 3, AR173: 3, AR254: 3, AR104: 3, AR197: 3, AR238: 3, AR286: 3, AR255: 2, AR266: 2, AR200: 2, AR296: 2, AR273: 2, AR297: 2, AR247: 2, AR237: 2, AR234: 2, AR267: 2, AR233: 2, AR242: 2, AR233: 2, AR247: 2, AR233: 2, AR247: 2, AR233: 2, AR2483: 2, AR228: 2, AR2483: 2, AR228: 2,	AR217: 2, AR211: 2, AR171: 2, AR229: 2, AR210: 1, AR055: 1, AR272: 1, AR216: 1, AR061: 1, AR289: 1 L0777: 3, H0624: 2, S0366: 2, L0779: 2, L0759: 2, L0604: 2, H0171: 1, H0149: 1, H0729: 1, H0741: 1, L0622: 1, L0623: 1, H0746: 1, H0213: 1, S0364: 1, H078: 1 and L0636: 1.	H0421: 1, L0118: 1, H0271: 1, S0448: 1, L0766: 1, S0216: 1, H0539: 1 and L0748: 1.	AR089: 28, AK310: 23, AR060: 18 L0439: 20, L0748: 16, L0438: 14, L0740: 12, L0731: 10, L0659: 9, L0803: 7, L0758: 7, L0759: 6, L0794: 5, H0050: 4, H0242: 4, L0771: 4, L0756: 4, L0779: 4, L0777: 4, L0757: 4, H0013: 3, H0599: 3, H0052: 3, T0010: 3, H0038: 3, L0770: 3, L0771: 3, L0775: 3, L0655: 3, L0809: 3, L0663: 3,
		2848	2849
		413 - 631	4310 - 4351
		251	252
		842176	839564
		HBMBD51	HBMBD73
		241	242

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	2850	2851	2852	2853	2854
	45 - 158	31 - 144	69 - 164	100 - 162	1815 - 1877
	253	254	255	256	257
	493916	637518	899809	821318	856461
	HBMBE33	HBMBM17	HBMCL59	нвмсм96	HBMCQ74
	243	244	245	246	247

	106165, 222900, 276902
	3q25 0740: 0774: 0581: 0779: 0779: 0779: 0779: 0779: 0779: 0779: 0574: 0014: 0036:
1, H0250: 1, S0280: 1, H0122: 1, H0004: 1, L0105: 1, H0318: 1, H00421: 1, H0196: 1, H0251: 1, H00421: 1, H0046: 1, H0196: 1, H00471: 1, H0046: 1, H0049: 1, L0471: 1, H0083: 1, H0015: 1, H0057: 1, H0015: 1, H0054: 1, H0057: 1, H0052: 1, H0053: 1, H0054: 1, H00640: 1, H00640: 1, H00640: 1, H00640: 1, H00640: 1, H00640: 1, L0774: 1, L0794: 1, L0803: 1, L0774: 1, L0657: 1, L0657: 1, L0657: 1, L0659: 1, L0664: 1, S0052: 1, L0666: 1, L0664: 1, S0052: 1, L0783: 1, H00435: 1, H0044: 1, H0365: 1, L0777: 1, S0028: 1, L0777: 1, S0028: 1, L0777: 1, S0028: 1, L0777: 1, S0050: 1, L0777: 1, S0050: 1, L0777: 1, S0050: 1, L0777: 1, S0050: 1, L0799: 1, H0057: 1, H00	AR060: 4, AR316: 4, AR089: 4 L0731: 17, L0766: 15, H0521: 15, H0457: 13, L0740: 9, H0486: 8, S0358: 6, S0474: 6, L0747: 6, H0423: 6, H0581: 5, L0754: 5, H0638: 4, L0779: 4, H0445: 4, S0134: 3, S0045: 3, S0374: 3, L0757: 3, L0485: 3, H0422: 3, S0040: 2, H0583: 2, S0360: 2, H0586: 2, H0574: 2, H0575: 2, H0516: 2, S0036: 2, H0575: 2, L0369: 2, H0500: 2, H0500: 2, H0500: 2, L0369: 2, H0500: 2, H0500: 2, H0500: 2, H0500: 2, H0500: 2, L0369: 2, H0500: 2, L0369: 2, H0500: 2, L0369: 2, L0360: 2, L03
	7 9 9 9 9 9 9 9 7 7 7 7
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	258 1815
	HBMCQ74 864382
	248

		1pter-p31.2	
2, L0764: 2, L0665: 2, L0765: 2, L07665: 2, L0665: 2, L0665: 2, L0665: 2, L0668: 2, L0668: 2, L0668: 2, L0750: 2, L07628: 1, H0171: 1, H0650: 1, H0650: 1, H0459: 1, H0650: 1, H0459: 1, H0650: 1, H0459: 1, H0650: 1, H0459: 1, H0650: 1, H0650: 1, H0650: 1, H0650: 1, H0650: 1, H0650: 1, H0660: 1, L0660: 1, L	1, S0026: 1, E0746: 1, E0749: 1, L0756: 1, L0777: 1, S0260: 1, L0590: 1, L0591: 1, H0667: 1 and L0697: 1.	AR089: 15, AR316: 13,	AR060: 12 L0748: 16, L0758: 10,
	1 1 1 1		Gin-49 to His-57.
		2856 G	5
		184 - 432	
		259	
		824053	
		HBMCT40	
		249	

																																					
: 7, L0747: 7,	: 6, L0776: 6,	. 4. 1.0757: 4.	: 3, H0494: 3,	: 3, L0665: 3,	3: 2, S0358: 2,	: 2, H0421: 2,	7: 2, T0010: 2,	3: 2, H0553: 2,): 2, S0142: 2,	: 2, L0768: 2,	L0803: 2, L0653: 2,	L0664: 2, S0374: 2,	: 2, L0759: 2,	: 2, L0485: 2,	: 1, S0114: 1,	I: 1, H0662: 1,	: 1, H0675: 1,	: 1, H0261: 1,	: 1, H0586: 1,	3: 1, H0069: 1,	7: 1, H0098: 1,	: 1, S0049: 1,	7: 1, H0545: 1,	: 1, H0510: 1,	3: 1, H0252: 1,	: 1, H0124: 1,): 1, H0280: 1,	: 1, S0144: 1,	: 1, H0529: 1,	Ę,	į,	: 1, L0771: 1,	L0533: 1, L0775: 1,	L0379: 1, L0657: 1,	: 1, H0593: 1,	: 1, H0670: 1,	, S03/8: 1, S0332: 1, . H0696: 1, H0436: 1.
L0754: 9, L0751: 7, L0747: 7,	10769: 6, L0806: 6, L0776: 6, 10430: 6, 10740: 5, 10500: 4	1.0662: 4.1.0774: 4.1.0757: 4.	H0506: 4, S0418: 3, H0494: 3,	L0659: 3, L0666: 3, L0665: 3,	H0521: 3, H0663: 2, S0358: 2,	L0717: 2, S0474: 2, H0421: 2,	H0052: 2, H0457: 2, T0010: 2	H0416: 2, H0188: 2, H0553: 2	H0135: 2, H0560: 2, S0142: 2,	L0369: 2, L0764: 2, L0768: 2,	L0766: 2, L0803		L0750: 2, L0731: 2, L0759: 2,	S0031: 2, L0596: 2, L0485: 2,	L0604: 2, S6024: 1, S0114: 1,	H0255: 1, H0661: 1, H0662:	H0638: 1, S0420: 1, H0675: 1.	S0132: 1, H0619: 1, H0261:	S0222: 1, H0455: 1, H0586: 1	H0587: 1, H0333: 1, H0069:	H0427: 1, H0097: 1, H0098:	T0082: 1, H0706: 1, S0049:	H0196: 1, H0597: 1, H0545:	H0024: 1, H0051: 1, H0510:	H0179: 1, H0288: 1, H0252:	H0615: 1, S0364: 1, H0124: 1,	H0163: 1, H0090: 1, H0280:	L0065: 1, H0647: 1, S0144: 1,	S0344: 1, L0598: 1, H0529: 1,			L0761: 1, L0646: 1,		L0523: 1, L0379	L0783: 1, H0144: 1, H0593:		H0648: 1, S03/8: 1 H0522: 1, H0696: 1
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	193300, 193300, 227646, 601154			
	3p25.3-3p24.1	4		
S3012: 1, S0028: 1, L0744: 1, L0779: 1, L0752: 1, S0394: 1, L0588: 1, L0608: 1, L0601: 1, H0667: 1, S0192: 1, S0276: 1 and S0196: 1.	H0421: 1	L0754: 4, L0766: 3, L0803: 3, H0648: 3, L0758: 3, L0800: 2, L0775: 2, L0666: 2, L0747: 2, L0779: 2, S0116: 1, S0358: 1, H0574: 1, L0021: 1, L0022: 1, L0667: 1, L0764: 1, L0768: 1, L0647: 1, L0647: 1, L0667: 1, L0777: 1, and 10599: 1	S0116: 1	L0439: 12, L0731: 11, L0754: 10, L0438: 8, L0750: 8, L0766: 7, L0752: 6, L0758: 6, L0759: 5, S0212: 4, H0266: 4, L0774: 4, H0556: 3, S0046: 3, L0775: 3, L0659: 3, L0779: 3, L0588: 3, S0420: 2, S0360: 2, H0333: 2, H0628: 2, T0010: 2, L0769: 2, L0772: 2, L0646: 2, L0773: 2, L0649: 2, L0740: 2, L0773: 2, L0659: 2, L0740: 2, L0769: 2, L0772: 2, L0606: 2, L0773: 2, L0655: 2, L0740: 2, L0773: 2, L0658: 2, L0740: 2, L0766: 2, L0755: 2, L0777: 2, L0769: 2, L0775: 2, L0606: 2, L0766: 2, L0755: 2, L0740: 1, S0040: 1, S0114: 1, T0049: 1, H0663: 1, S0118: 1, S0354: 1, H0663: 1, S0354:
	Asp-85 to Arg-92, Ala-107 to Glu-116.			
	2857	2858	2859	2860
	505 - 975	193 - 288	172 - 225	44 - 58
	260	261	262	563
	837927	787680	843389	886609
	HBMDM08	HBMSN62	HBMSO30	
	250	251	252	253

	141750, 141800, 141800, 141800, 141800, 141850, 141850, 141850, 141850, 156850, 156850, 191092, 600140, 600273, 601785
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	Arg-31 to Cys-38, Thr-69 to Thr-76, Pro-94 to Glu-100, Ile-114 to Glu-120.
	2861
	270 - 650
	264
,	701970
	HBMUD59
	254

				20, AR273:
				18, AR313:
				18, AR186:
				17, AR266:
				16, AR185:
				14, AR175: 1
				AR256: 13, AR061: 13,
				AR219: 12, AR253: 12,
				AR231: 12, AR293: 12,
				AR248: 12, AR183: 12,
	-	-		
				AR269: 11, AR233: 11,
				AR291: 11, AR296: 11,
				AR289: 11, AR259: 11,
	-	-	-	AR229: 11, AR258: 10,
				AR298: 10, AR226: 10,
,				10, AR249:
				8, AR182:
				AR267: 7, AR184: 6,
				AR179: 6
				L0794: 6, L0803: 6, L0663:
	-			5, L0758: 5, H0617: 4, L0731:
	-			4, L0804: 3, L0806: 3, L0665:
				[3, L0439: 3, L0747: 3, H0556:
				2, S0360: 2, H0581: 2, H0424:
				2, L0769: 2, L0662: 2, L0768:
-				2, L0809: 2, L0791: 2, H0684:
				2, S0380: 2, L0743: 2, L0744:
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				1, S0418: 1, S0354: 1, S0300:
				1, H0052: 1, H0546: 1, H0545:
				1, H0012: 1, H0275: 1, H0687:
				1, H0030: 1, H0644: 1, H0182:

19951083.191201

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	Tyr-5 to Leu-10. 3, 3, 2, 2, 2, 2, 2, 2, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,	A. A
	2862	2863
	249 - 368	180 - 254
	265	266
	566764	580808
	HBMUI10	HBMUJ48
	255	256

D9950083.091201

		,					12
1, H0428: 1, H0169: 1, L0455: 1, T0042: 1, S0002: 1, L0638: 1, L0772: 1, L0771: 1, L0651: 1, L0663: 1, L0665: 1, H0547: 1, H0521: 1, S0027: 1 and L0757: 1.	AR089: 18, AR316: 14, AR060: 11 S0116: 1 and L0748: 1.	L0803: 2, L0526: 2, S0116: 1, S6028: 1, L0627: 1, L0777: 1, L0759: 1, L0608: 1, S0026: 1 and S0462: 1.	L0803: 2, L0526: 2, S0116: 1, S6028: 1, L0627: 1, L0777: 1, L0759: 1, L0608: 1, S0026: 1 and S0462: 1.	H0590: 2, S0002: 2, S0126: 2, H0521: 2, H0265: 1, T0049: 1, S0116: 1, H0638: 1, S0360: 1, H0004: 1, H0030: 1, T0041: 1, S0344: 1, L0794: 1, L0766: 1, L0806: 1, L0805: 1, L0543: 1, L0749: 1, H0136: 1 and H0423: 1.	L0587: 2, H0170: 1, S0116: 1, S0442: 1, H0431: 1, H0574: 1, H0318: 1 and H0615: 1.	S0116: 1	AR089: 10, AR316: 10, AR060: 10 L0740: 4, H0616: 3, H0056: 3, L0803: 3, L0659: 3, L0666: 3, L0758: 3, L0005: 2, L0717: 2, H0032: 2, L0455: 2, L0598: 2, L0800: 2, L0771: 2, L0766: 2, L0663: 2, L0665: 2, L0438: 2, L0748: 2, L0759: 2, H0686: 1, H0583: 1, S0116: 1, H0638:
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	2864	2865	2866	2867	2868	5869	2870
	135 - 206	179 - 295	382 - 498	112 - 168	82 - 204	921 - 61	301 - 336
	267	268	269	270	271	272	273
	647594	609348	847789	523713	701971	872553	834497
	HBMUR39	HBMVF65	HBMVF65	нвммсз9	HBMWJ92	HBMWS52	HBMXE34
	257	258	259	260	261	262	263

				180297, 230450, 263200, 601690
1, \$0348: 1, \$0354: 1, \$0476: 1, \$0222: 1, H0431: 1, L0021: 1, \$0202: 1, H0431: 1, L0021: 1, H0545: 1, \$0050: 1, \$0051: 1, H0688: 1, H0428: 1, H0553: 1, H0124: 1, H0090: 1, H0591: 1, H0038: 1, H0488: 1, \$0448: 1, \$0440: 1, \$0002: 1, L0770: 1, L0646: 1, L0764: 1, L0649: 1, L0804: 1, L0653: 1, L0776: 1, L0804: 1, L0809: 1, \$0374: 1, H0519: 1, H0435: 1, H0521: 1, H0696: 1, L0751: 1, L0747: 1, L0780: 1, \$0260: 1, \$0434: 1, L0595: 1, H0665: 1, \$0242: 1, L0698: 1 and H0506: 1.	S0116: 1	AR060: 7, AR316: 5, AR089: 3 S0116: 1	L0748: 6, S0114: 1, S0116: 1, H0431: 1, H0313: 1, L0740: 1, L0749: 1, L0758: 1 and S0260: 1.	S0410: 9, L0803: 5, L0809: 4, L0756: 3, L0759: 3, H0171: 2, S0116: 2, H0586: 2, S0003: 2, L0769: 2, L0662: 2, L0805: 2, L0653: 2, L0777: 2, L0749: 2, L0779: 2, L0777: 2, L0752: 2, L0777: 2, L0752: 1, H0664: 1, H0604: 1, H0604: 1, H0604: 1, H0604: 1, H0604: 1, H0604: 1, H0606: 1, H0606: 1, H0606: 1, H0606: 1, L0763: 1, L0606: 1, L0763: 1, L0606: 1, L0763: 1, L0766: 1, L0768: 1, L0768: 1, L0768: 1, L0768: 1, L0768: 1, L0388: 1, L0804: 1, L0774:
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	2871	2872	2873	2874
	261 - 335	71 - 151	20 - 133	16 - 135
	274	275	276	277
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·	264	265	266	267

ISSECTES OSTECT

120550, 120570, 120575, 121800, 130500, 133200, 138140, 171760, 171760, 171760, 600975	
1p36.11-p34.2	16
1, L0775: 1, L0806: 1, L0606: 1, L0659: 1, L0788: 1, L5286: 1, L0666: 1, L0663: 1, L0665: 1, L0665: 1, L0665: 1, L0748: 1, L0731: 1, S0434: 1 and H0422: 1. AR060: 5, AR316: 4, AR089: 3 H0441: 3, L0806: 2, L0663: 2, L0750: 2, H0188: 1, H0688: 1, H0428: 1, L0749: 1, S0028: 1, L0749: 1, S0028: 1, L0749: 1, and L0758: 1.	H0271: 10, H0556: 7, L0731: 7, H0494: 5, L0766: 5, L0777: 5, H0622: 4, L0805: 4, L0741: 4, L0743: 4, L0744: 4, L0749: 4, H0619: 3, H0639: 3, H0619: 3, H0629: 2, L0740: 3, H0629: 2, L0769: 2, L0667: 2, L0667: 2, L0667: 2, L0667: 2, L0776: 2, L0777: 2, L0778: 2, L0776: 2, L0777: 2, L0778: 2, L0776: 2, L0777: 2, L0778: 2, L0776: 2, L0777: 1, R0642: 1, H0638: 1, H0642: 1, H0649: 1, H0659: 1, H0657: 1, H0659: 1, H0657: 1, H0657
	Gly-62 to Gly-69, Pro-96 to Asp-102.
2875	2876
	2711 - 3157
278	279
637524	843727
HBNAE74	HBNAX16
268	269

TOSTOOT EBOOTEST

1, H0050: 1, H0099: 1, H0031: 1, H0617: 1, H0616: 1, T0042: 1, H0560: 1, H0560: 1, L0378: 1, L0378: 1, H0690: 1, H06	i, 10156: : 1.	4, , L0764: 3, S0360: 2, H0624: 1, S0358: 1, H0052: 1, S0344: 1, S0344: 1, L0646: 1, L0646: 1, L0664: 1, S0330:
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	2877	2878
	519 - 542	338 - 370
	280	281
	840687	852382
	HBNAZ35	HBODK40
	270	271

	· · · · · · · · · · · · · · · · · · ·	
	203800	
	2p14-q14.3	
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	2879	7880
	303 - 329 28	74 - 121 28
	782	283
	76 866420	99 637525
	HBODV76	HBPAD89
	272	273

			3, L0750: 3, H0445: 3, S0434: 3, S0436: 3, H0422: 3, T0002:		
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			2, H0575: 2, H0618: 2, H0620:		
			2, H0373: 2, H0083: 2, H0266:		
			2, H0271: 2, H0687: 2, S0250:		
			2, H0615: 2, L0194: 2, H0553:		
			2, H0090: 2, H0038: 2, H0059:		
			2, H0100: 2, T0041: 2, H0561:		
			2, L0761: 2, L0372: 2, L0646:		
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			2 H0648: 2, S0378: 2, S3014:		
			2, 1100-10; 2, 500-10; 2, 500-11; 2, 500-20; 2, 1,0754; 2, 1,0753;		
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			1, S0001: 1, S0030: 1, H0255:		
			1, H0661: 1, H0663: 1, H0125:		
			1, S0354: 1, S0358: 1, S0468:		
			1, S0132: 1, S0476: 1, H0619:		
	· ·		1, H0411: 1, S0278: 1, H0613:		
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			[1, T0039: 1, L0021: 1, H0706:		
	•		1, S0010: 1, H0390: 1, S0049:		
			1, H0596: 1, H0597: 1, H0457:		
			1, H0041: 1, H0009: 1, N0006:		
			1, H0563: 1, H0566: 1, H0050:		
			1, H0023: 1, H0014: 1, S0362:		
			1, H0051: 1, S0048: 1, T0010:		
			1, S6028: 1, H0028: 1, S0314:	-	
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			Thr-49 to Gly-55, Ser-70 to Glu-77.	
	2881	2882	2883	2884
	119 - 175	292 - 315	741 - 974	208 - 321
	284	285	286	287
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	HBPAF39	HBQAC45	HBQAC72	НВQАЕ37
	274	275	276	277

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170650, 600900	133701, 168500,	171650, 176930,	176930, 600623,	600811, 600958																																		
	11p11.2-p11.12																																					
	H0253: 8, L0439: 8, L0769:	7, H0618: 6, L0758: 6, H0052:	5, L0749: 5, H0617: 4, H0135:	4, L0766: 4, S0406: 4, S0001:	3, H0255; 3, S0410; 3, H0619;	3, S0422: 3, L0775: 3, L0378:	3, H0547; 3, H0521; 3, L0742;	3, L0750: 3, L0755: 3, L0757:	3 S0434·3 L0605·3 H0381·	2 HOA10: 2 HO241: 3 SO420:	2, IIO419: 2, IIO541: 2, 30420:	2, HU33U: 2, HU438: 2, HU399:	2, H0318: 2, H0046: 2, H0050:	2, H0012: 2, H0024: 2, S0050:	2, T0010: 2, L0455: 2, H0412:	2, H0413: 2, H0494: 2, L0772:	2, L0645: 2, L0764: 2, L0771:	2, L0438: 2, H0520: 2, H0519:	2, H0134: 2, 1,0741: 2, 1,0748:	2. L0751: 2. L0747: 2. L0777:	2, L0759: 2, H0445: 2, L0596:	2, L0603: 2, L0411: 1, H0556:	1, S0116: 1, H0125: 1, S0418:	1, S0354: 1, S0360: 1, H0729:	1, H0730: 1, H0722: 1, H0728:	_	1, H0549: 1, H0370: 1, H0392:	1, H0613: 1, H0013: 1, H0427:	1, H0575: 1, T0082: 1, H0706:	1, H0036: 1, H0421: 1, S0049:	1, H0194: 1, H0085: 1, H0231:	1, L0041: 1, H0041: 1, H0009:	1, H0123: 1, H0620: 1, H0199:	1, H0246: 1, H0014: 1, L0163:	1, H0594: 1, S6028: 1, H0266:	1, H0188: 1, H0687: 1, H0288:	1, H0033: 1, H0181: 1, S0364:	1, S0366: 1, S0036: 1, H0038:
	2885																																					
	1256 - 1444																																			·		
	288																																					
	848683																																					
	HBSAJ63																																					
	278																																					

DGGSOLESOLESOL

	133701, 168500, 171650, 176930, 176930, 600623, 600811, 600958
	11p11.2-p11.12
1, H0616: 1, H0264: 1, H0268: 1, H01017: 1, S0038: 1, H0100: 1, L0351: 1, L0435: 1, T0041: 1, T0042: 1, S0448: 1, S0142: 1, S0002: 1, H0529: 1, L0796: 1, L0763: 1, L0768: 1, L0768: 1, L0774: 1, L0648: 1, L0768: 1, L0776: 1, L0559: 1, L0550: 1, L0550: 1, L0550: 1, L0776: 1, L0783: 1, L0793: 1, H0144: 1, H0539: 1, S0152: 1, H0650: 1, S0330: 1, H0539: 1, S0330: 1, S0028: 1, L0743: 1, S0152: 1, L0752: 1, H0644: 1, S0330: 1, S0028: 1, L0743: 1, L0779: 1, L0752: 1, H0644: 1, H0555: 1, L0752: 1, H0644: 1, H0555: 1, L0752: 1, H0644: 1, S0330: 1, S0028: 1, L0743: 1, L0779: 1, L0752: 1, H0644: 1, S0436: 1, L0752: 1, H0543: 1, H0543: 1, H0543: 1, H0550: 1, S0458: 1 and H0506: 1.	H0253: 8, L0439: 8, L0769: 7, H0618: 6, L0758: 6, H0052: 5, L0749: 5, H0617: 4, H0135:: 4, L0766: 4, S0406: 4, S0001: 3, H0255: 3, S0410: 3, H0619: 3, S0422: 3, L0775: 3, L0742: 3, L0750: 3, L0775: 3, L0742: 3, L0750: 3, L0755: 3, L0742: 3, S0434: 3, L0651: 3, L0742: 2, H0419: 2, H0341: 2, S0420: 2, H0318: 2, H0448: 2, H0599: 2, H0413: 2, H0449: 2, L0772: 2, L0645: 2, L0666: 2, L0665: 2, L0662: 2, L0666: 2, L0665: 2, L0771: 2, L0741: 2, L0751: 2, L0751: 2, L0751: 2, L0751: 2, L0596: 2, L0596
	7886 2
	183 - 371
	835454 289
	HBSAJ63 835
	279

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2, L0603: 2, L0411: 1, H0556: 1, S0114: 1, S0218: 1, H0656: 1, S0114: 1, S0218: 1, H0656: 1, S0116: 1, H0125: 1, S0418: 1, H0730: 1, H0722: 1, H0728: 1, H0733: 1, L0717: 1, S0278: 1, H0613: 1, H0013: 1, H0427: 1, H063: 1, H0031: 1, H0427: 1, H0613: 1, H0041: 1, H0099: 1, H0194: 1, H0041: 1, H0099: 1, H0194: 1, H0041: 1, H0266: 1, H0194: 1, H0041: 1, H0266: 1, H0194: 1, H0041: 1, L0163: 1, H017: 1, S0038: 1, H0266: 1, L076: 1, L0374: 1, L0766: 1, L076: 1, L0374: 1, L0776: 1, L076: 1, L0374: 1, L0776: 1, L076: 1, L0793: 1, H0144: 1, H0690: 1, H0660: 1, S0330: 1, H0522: 1, H0644: 1, S0436: 1, L0752: 1, H0444: 1, S0436: 1, L0752: 1, H0560: 1, S0330: 1, L0752: 1, H0560: 1, S0330: 1, L0752: 1, H0444: 1, S0436: 1, L0752: 1,	AR089: 9, AR316: 9, AR060: 9 L0761: 4, L0766: 4, L0747:
	2887
	833 - 961
	290
	839802
	HBSDD24
	280

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			2pter-p25.1
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2 2 2 2 2 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1	P	7	His-25 to Phe-30.
	2888	2889	2890
	182 - 328	215 - 250	1097 - 1219
	291	292	293
	800765	836513	815571
	HBWBD25	HBXAS93	HBXAT27
	281	282	283

				103850, 114835, 121360, 217800, 218030
				16q22
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	7	Met-I to Ala-7.		Arg-24 to Gly-29, Gly-43 to Arg-49.
	2891	2892	2893	2894
	152 - 262	87 - 173	85 - 225	2163 - 2522
	294	295	296	297
	815650	847000	823353	812527
	HBXAW57	HBXBI29	HBXBM24	HBXBM78
	284	285	286	287

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1, H0271: 1, S0038: 1, S0152: 1, H0521: 1 and H0522: 1.	L0439: 5, L0438: 3, L0769: 2, L0759: 2, S0001: 1, S0282: 1, S0300: 1, S0222: 1, H0438: 1, S0010: 1, S0049: 1, H0052: 1, H0572: 1, T0010: 1, S0038: 1, H0521: 1, L0756: 1 and L0366: 1.	S0038: 1	AR089: 14, AR316: 9, AR060: 5 H0031: 3, L0741: 3, L0757: 2, L0032: 1, H0123: 1, L0455: 1, H0135: 1, S0038: 1, H0100: 1, L0659: 1, L0792: 1, L0665: 1, S0380: 1 and L0591: 1.	AR089: 11, AR316: 7, AR060: 4 S0222: 3, H0100: 2, L0438: 2, L0759: 2, L0592: 2, L0593: 2, S0114: 1, S0001: 1, H0208: 1, H0123: 1, H0012: 1, H0620: 1, S6028: 1, H0328: 1, H0413: 1, S0038: 1, T0042: 1, H0547: 1, L0439: 1 and L0608: 1.	L0758: 9, L0752: 8, L0731: 6, L0748: 5, L0740: 5, L0747: 5, H0014: 4, L0774: 4, L0754: 4, L0756: 4, L0755: 4, S0358: 3, S0360: 3, H0083: 3, H0032: 3, H0059: 3, L0770: 3, L0766: 3, L0757: 3, H0171: 2, H0265: 2, H0657: 2, H0156: 2, H0428: 2, L0769: 2, L0806: 2, L0655: 2, H0659: 2, L0866: 2, L0655: 2, H0659: 2, L0866: 2, L0655: 2, H0659: 2, S0152: 2, H0521: 2, H0659: 2, S0152: 2, L0439:
	Met-1 to Cys-7, Leu-21 to Arg-44, Ala-53 to Ser-60.			2 2 2 1 1 1	Thr-4 to Arg-10, Pro-42 to Ser-47. 3 3 2 2 2 2 2
	2895	2896	2897	2898	2899
	165 - 362	349 - 363	260 - 313	366 - 479	231 - 398
	298	299	300	301	302
	860439	801608	628501	799513	589516
	HBXCD59	HBXCE43		HBXCM52	нвхсооз
	288	289	290	291	292

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6. 8. 7. 9. 9. 9. 9. 9. 9. 9. 9. 9. 9. 9. 9. 9.	7: 19p13.3 9: 00: 22: 55: 15: 15: 16: 17: 17: 19p13.3	88: 05: 72:
2, L0588: 2, S0026: 2, H06248: 1, S0402: 1, T0049: 1, S0420: 1, L0005: 1, S0354: 1, H0489: 1, S0045: 1, S0045: 1, S0046: 1, S0046: 1, S0045: 1, S0046: 1, S0036: 1, H00592: 1, H0013: 1, S0278: 1, H0577: 1, H0013: 1, H0599: 1, H0179: 1, H0647: 1, H019: 1, H0568: 1, H0649: 1, S0038: 1, H0649: 1, S0150: 1, H0641: 1, S0038: 1, H0494: 1, L0772: 1, L0764: 1, L0773: 1, L0768: 1, L0764: 1, L0773: 1, L0768: 1, H0506: 1 and H0345: 1	L0439: 18, H0441: 8, L0157: 7, S0007: 3, H0052: 3, L0769: 3, S0001: 2, S0222: 2, H0100: 2, L0805: 2, L0756: 2, S0282: 1, S6026: 1, H0261: 1, H0455: 1, H0194: 1, H0009: 1, H0201: 1, S0388: 1, T0010: 1, S6028: 1, H0087: 1, S0388: 1, S0388: 1, L0776: 1, L0779: 1, L0776: 1, L0776: 1, L0741: 1, L0742: 1, L0786: 1 and L0592: 1.	L0438: 4, L0439: 3, L0779: 3, L0785: 2, L0471: 2, S0038: 2, L0521: 2, S0116: 1, S0360: 1, H0637: 1, S0222: 1, H0486: 1, H0013: 1, H0004: 1, H0052:
	Gly-17 to Ala-22, Tro-24 to Pro-33. 7, 7, 1, 1, 1, 1, 1, 1, 1, 1,	3, 2, 3, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
	2900 G	2901
	158 - 259	1040 - 1066
	303	304
	731861	873460
	HBXCR15	HBXDL52
	293	294

		13
1, H0012: 1, H0038: 1, H0616: 1, L0803: 1, L0789: 1, S0053: 1, L0731: 1, L0758: 1, L0599: 1 and L0594: 1.	L0438: 4, L0439: 3, L0779: 3, L0785: 2, L0471: 2, S0038: 2, L0521: 2, S0116: 1, S0360: 1, H0637: 1, S0222: 1, H0486: 1, H0013: 1, H0004: 1, H0052: 1, H0012: 1, H0038: 1, H0616: 1, L0803: 1, L0789: 1, S0053: 1, L0731: 1, L0758: 1, L0599: 1 and L0594: 1.	AR175: 16, AR270: 14, AR284: 11, AR289: 11, AR295: 11, AR290: 10, AR295: 11, AR290: 10, AR293: 10, AR268: 10, AR293: 10, AR296: 10, AR293: 10, AR296: 10, AR292: 9, AR184: 8, AR296: 7, AR298: 7, AR286: 7, AR298: 7, AR219: 7, AR298: 7, AR240: 6, AR186: 6, AR289: 6, AR089: 6, AR289: 5, AR291: 5, AR291: 5, AR294: 5, AR291: 5, AR294: 5, AR291: 4, AR192: 4, AR296: 3, AR298: 3, AR295: 3, AR298: 3, AR275: 3, AR858: 3, AR295: 3, AR298: 3, AR295: 3, AR298: 3, AR295: 3, AR298: 3,
	2902	2903
	1040 - 1066	67 - 141
	305	306
	889328	566765
	HBXDL52	HBXDN08
	295	296

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AR055: 3. AR229: 2.				` '			_	· -	1 0777 14 1 0439 11	748. 10 T 0747. 10 T 0754.	LU/48: 10, LU/4/: 10, LU/54:	8, LU8U3: /, LU/31: /, LU/31:	7, S0212: 6, H0628: 5, L0794:	5, L0809: 5, L0741: 5, L0750:	5, L0759: 5, S0007: 4, L0771:	4, L0662: 4, H0144: 4, L0749:	4, L0756: 4, S0358: 3, H0575:	3, S0038: 3, L0776: 3, L0517:	3, H0670: 3, S0380: 3, S0028:	3, 1.0744; 3, 1.0758; 3, 1.0592;	3, H0717: 2, H0663: 2, S0418:	2, S0420: 2, H0619: 2, S0222:	2, S0010: 2, H0545: 2, H0009:	2, H0266: 2, H0615: 2, H0412:	2, S0440: 2, L0770: 2, L0769:	2, L0637: 2, L0764: 2, L0768:	2, L0775: 2, L0526: 2, L0783:	2, L0790: 2, L0438: 2, L0352:	2, H0547: 2, H0658: 2, H0696:	2, L0742: 2, L0752: 2, L0594:	2, L0603: 2, H0624: 1, H0170:	S0001: 1, S0400: 1, S0442:	S0354: 1, S0376: 1, S0408:	H0393: 1, H0549: 1, H0438:	, H0333: 1, H0574: 1, H0270:	1, L0477: 1, T0114: 1, H0196:	1, H0052: 1, H0251: 1, L0738:	1, H0327: 1, H0567: 1, H0123:	1, L0471: 1, H0011: 1, H0012:	H0620: 1 H0083: 1 H0099:
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		Gly-25 to Glu-34.			Pro-87 to Glu-96, Thr-98 to Thr-103.
	2904	2905		2906	2907
	187 - 297	411 - 536		15 - 116	124 - 495
	307	308		309	310
	840021	842901		838824	610452
	HBXDN65	HBXFA04		HBXFE64	
	297	298		299	300

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1, S0003: 1, H0039: 1, T0023: 1, L0796: 1, L0766: 1, L0653: 1, L0665: 1, L0666: 1, L0666: 1, L0666: 1, H0547: 1, H0689: 1, H0648: 1, H0577: 1, L0759: 1 and H0543: 1.	H0438: 1 and L0759: 1. L0439: 9, T0010: 6, L0592: 6, L0500: 3, L0759: 3, L0794: 2, L0442: 1, S0282: 1, S0007: 1, S0220: 1, H0438: 1, H0434: 1, H0562: 1, H0327: 1, H0009: 1, H0568: 1, L0776: 1, L0351: 1, L0768: 1, L0741: 1, L0786: 1 and S0106: 1.	L0769: 10, L0439: 7, S0049: 2, H0052: 2, S0051: 2, L0742: 2, S0110: 1, L0617: 1, S0300: 1, H0441: 1, H0438: 1, H0485: 1, H0748: 1, H0778: 1, L0770: 1, L0639: 1, L0771: 1, L0768: 1, L074: 1, L078: 1, S0053: 1, H0539: 1, L0749: 1, L0758: 1, L0759: 1, L0758: 1, L0759: 1, L0758: 1, L0759:	AR039: 7, AR198: 5, AR247: 4, AR263: 4, AR265: 4, AR310: 3, AR204: 3, AR294: 2, AR249: 2, AR292: 2, AR296: 2, AR293: 2, AR192: 2, AR240: 2, AR205: 2, AR033: 2,
	& & & & & & & & & & & & & & & & & & &	Asp-39 to Ala-50, Lys-80 to Cys-85. Asp-39 to Ala-50, Lys-80 to Cys-85.	
	2910	2911	2912
	20 - 70 125 - 670	52 - 414	80 - 190
	313	314	315
	847001 847001	1310891	901845
	HBXFW01	HBXGE12	HBXGL91
	302	304	305

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			601843
			19p11-p12
AR274: 2, AR096: 2, AR295: 2, AR314: 2, AR312: 2, AR183: 2, AR284: 2, AR285: 2, AR269: 1, AR104: 1, AR298: 1, AR286: 1, AR281: 1, AR282: 1, AR182: 1, AR282: 1, AR244: 1, AR206: 1, AR299: 1, AR277: 1, AR299: 1, AR290: 1, AR186: 1 H0438: 1 and H0615: 1.	S0354: 1, S0222: 1, H0431: 1, H0438: 1, H0051: 1, S0426: 1 and H0529: 1.	H0339: 1 and S0188: 1.	L0761: 3, L0794: 3, H0271: 2, H0252: 2, L0769: 2, L0771: 2, L0766: 2, L0809: 2, H0521: 2, L0752: 2, L0747: 2, L0756: 2, L0752: 2, L0747: 2, L0756: 1, S0376: 1, H0609: 1, H0486: 1, H0439: 1, S0051: 1, H0688: 1, H0634: 1, H0413: 1, L0763: 1, L0770: 1, L0800: 1, L0800: 1, S0031: 1 and L0595: 1, H0125: 1 H0125: 1 L0758: 13, L0756: 3, L0752: 3, L0758: 13, L0758: 1, H0020: 1, H0309: 1, H0616: 1, H0202: 1, H0309: 1, L0766: 1, L0774: 1, L0369: 1, L0776: 1, L0369: 1, L0776: 1, H0547: 1, H0547: 1, H0690: 1, H0647: 1, H0647: 1, H06409: 1, H0690: 1, S0152: 1, H04499: 1, H0690: 1, S0152: 1, H0479: 1, H0690: 1, S0152: 1, H04499: 1, H0690: 1, S0152: 1, H0479: 1, H0690: 1, S0152:
		Gln-18 to Glu-24, Ser-39 to Lys-44.	Arg-31 to Met-36.
	2913	2914	2915 2916 2917 2918
·	268 - 315	260 - 406	9 - 122 76 - 198 30 - 140 93 - 212
	316	317	318 319 320 321
	821320	668231	866421 846467 850785 847002
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	306	307	308 309 310 311

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1, L0743: 1, L0439: 1, L0750: 1, L0777: 1, L0755: 1, H0445: 1 and L0588: 1.	AR183: 16, AR177: 11, AR290: 5, AR285: 4, AR284: 4, AR060: 6, AR284: 4, AR089: 4, AR294: 4, AR316: 4, AR299: 3, AR247: 3, AR179: 3, AR182: 3, AR179: 3, AR182: 3, AR240: 2, AR270: 2, AR299: 2, AR213: 2, AR299: 2, AR213: 2, AR299: 1, AR281: 1, AR299: 1, AR282: 1, AR299: 1, AR282: 1, AR299: 1, AR282: 1, AR299: 1, AR061: 1, AR299: 1, AR282: 1, AR299: 1, L0679: 1, L0759: 1, AR299: 1, L0739: 1, L0759: 1, AR299: 1, L0753: 1, L0759: 1, AR299: 1, AR29	AR183: 16, AR177: 11, AR175: 7, AR060: 6, AR290: 5, AR285: 4,
	Ala-85 to Thr-90, Glu-149 to Gly-155, Pro-157 to Asn-166.	A A A
	2919	2920
	342 - 890	427 - 855
	322	323
	806565	894773
	HCDAA24	HCDAA24
	312	313

	300075, 300077, 301200, 302350, 302801, 305435, 306000, 306000, 306100, 307800, 308800, 309510, 311200, 311770, 312040, 312170,
	Xp22.1-p22.2
AR284: 4, AR089: 4, AR294: 4, AR089: 4, AR294: 4, AR316: 4, AR269: 4, AR295: 3, AR289: 3, AR247: 3, AR179: 3, AR182: 3, AR299: 2, AR313: 2, AR299: 2, AR313: 2, AR240: 2, AR313: 2, AR266: 2, AR231: 2, AR266: 2, AR231: 2, AR266: 1, AR282: 1, AR291: 1, AR086: 1, AR292: 1, AR282: 1, AR292: 1, AR096: 1, AR292: 1, AR296: 1, AR292: 1, AR292: 1, AR292: 1, AR296: 1, AR292: 1, AR292: 1, AR292: 1, AR296: 1, AR292: 1, AR293: 1, L0772: 1, L0659: 1, AR293: 1, L0773: 1 and AR293: 1, L073: 1 and AR293: 1, L073: 1 and	2, H0551: 1, L0438: 7, S0356: 4, H0251: 4, L0649: 3, L0803: 3, L0754: 3, H0156: 2, L0471: 2, H0531: 2, H0551: 2, L0771: 2, H0550: 1, H0341: 1, S0448: 1, S0448: 1, S0448: 1, S0300: 1, H0486: 1, H0266: 1, H0673: 1, H0486: 1, H0560: 1, H0673: 1, H0488: 1, H0560: 1, H0673: 1, H0673: 1, H0673: 1, H0560: 1, H0673: 1, H06
AR234: AR294: AR294: AR269: AR179: AR260: AR260: AR260: AR260: AR260: AR260: AR260: AR260: AR260: AR293: AR294: AR	2, 4, 4, 1, 1, 1, 80, 11, 80,
	2921
	12 - 119 29
	324
	7 722206
	HCDAF17
	314

D9950083.091201

	109565, 109565, 142640, 228960, 261515, 600044
2	3427
1, H0509: 1, L0369: 1, L0763: 1, L0662: 1, L0794: 1, L0659: 1, L0789: 1, L0663: 1, S0126: 1, H0435: 1, H0660: 1, H0539: 1, S0128: 1, S0152: 1, S0037: 1, S0028: 1, L0756: 1, L0755: 1, L0759: 1 and S0242: 1. AR089: 11, AR316: 10, AR060: 9 H0265: 1, H0556: 1, H0251: 1, L0777: 1 and L0731: 1.	H0694: 54, L0731: 23, L0747: 21, L0748: 12, L0758: 12, L0766: 10, L0769: 8, L0775: 8, H0599: 7, L0776: 7, L0749: 7, L0771: 6, L0770: 5, L0519: 5, S0374: 5, L0750: 5, L0757: 5, H0685: 4, S0007: 4, S0222: 4, L0623: 4, H0413: 4, S0144: 4, L0761: 4, H0659: 4, L0752: 4, H0657: 3, H0341: 3, S0278: 3, H0392: 3, H0024: 3, H0266: 3, H0031: 3, S0038: 3, H0266: 3, H0031: 3, S0038: 3, L0803: 3, L0805: 3, L0558: 3, L0759: 3, S0440: 1, L0755: 3, H0170: 2, H0171: 2, T0002: 2, H0049: 2, L0808: 2, S0116: 2, H0049: 2, S0132: 2, H0674: 2, H0040: 2, H0087: 2, S0438: 2, H0646: 2, S0344: 2, L0772: 2, L0764: 2, L0521: 2, L0768: 2, H0658: 2, S0004: 2, H0666: 2, H0134: 2, S0206: 2, L0581: 2, L0756: 2, S0031: 2, L0581: 2, H0653: 2, H0653: 2, H0653: 2,
Glu-40 to Thr-46.	
2922	2923
495 - 665	126 - 209
325	326
653066	566794
нсран02	HCDAP33
315	316

H0395: 1, H0686: 1, H0294: 1, S0212: 1, H0484: 1, L0785: 1, S0212: 1, H0484: 1, H0483: 1, H0662: 1, S0358: 1, S0444: 1, S0360: 1, S0360: 1, S0360: 1, H0453: 1, H0491: 1, H0492: 1, H0492: 1, H0492: 1, H0492: 1, H0492: 1, H0592: 1, H0698: 1, H0592: 1, H0698: 1, H0541: 1, H0698: 1, H0541: 1, H0698: 1, H0541: 1, H0698: 1, H0541: 1, H0698: 1, H0699:	S0114: 1, S0045: 1, H0318: 1 and H0251: 1.	AR089: 12, AR316: 12, AR060: 12
		Thr-70 to Ala-77.
	2924	2925
	102 - 224	136 - 870
	327	328
	654821	896667
	HCDAR40	HCDAS02
	317	318

	182600, 232700, 602086
	14q21.1-q21.3
L0439: 7, H0617: 6, H0181: 5, H0661: 3, L0769: 3, L0809: 3, L0751: 3, H0657: 2, S0408: 2, S0222: 2, H0618: 2, H0560: 2, L0764: 2, L0648: 2, L0665: 1, H0556: 1, S0409: 1, H0716: 1, H0656: 1, H0341: 1, H0484: 1, H0656: 1, H0580: 1, S0300: 1, H055: 1, H0580: 1, S0300: 1, H055: 1, H0580: 1, R0350: 1, H0675: 1, H0580: 1, H0251: 1, H0675: 1, H0569: 1, H0251: 1, H0614: 1, H0569: 1, H0267: 1, H0614: 1, H0569: 1, H0267: 1, H0614: 1, L0769: 1, H0640: 1, L0642: 1, L0643: 1, L0760: 1, L0642: 1, L0643: 1, L0760: 1, L0642: 1, L0643: 1, L0760: 1, L0642: 1, L0666: 1, L0789: 1, L0662: 1, L0666: 1, L0789: 1, L0791: 1, L0666: 1, L0663: 1, L0791: 1, L0666: 1, H0555: 1, S0028: 1, L078: 1, H0779: 1, S0042: 1, S0448: 1, L0779: 1, L078: 1, L0779: 1, L0779: 1, L078: 1, L0779: 1, L078: 1, L0779: 1, L078: 1, L0779: 1, L078: 1, H0779: 1, S0042: 1.	AR089: 9, AR316: 8, AR060: 8 L0748: 17, L0740: 7, L0777: 5, S0140: 4, H0457: 4, H0013: 3, H0615: 3, S0002: 3, L0763: 3, H0665: 3, L0439: 3, L0758: 3, H0662: 2, H0628: 2, L0766: 2, L0803: 2, L0809: 2, L0750: 2, L0756: 2, L0755: 2, L0731: 2, H0170: 1, S0001: 1, H0580: 1, H0411: 1, H0575: 1, T0082: 1, H0251: 1, T0115: 1, L0471:
V (V)(V)(V)	
	2926
	260 - 406
	329
	840076
	нсрве76
	319

COSTOCE COSTOCE

		20	
1, \$0003: 1, H0688: 1, H0031: 1, L0456: 1, H0591: 1, L0369: 1, L074: 1, L0749: 1, L0774: 1, L0375: 1, L0776: 1, H0591: 1, H0519: 1, H0689: 1, H0539: 1, S0380: 1, H0696: 1, L074: 1, L0779: 1, L0779: 1, L0779: 1, L0780: 1, L0759: 1, L0789: 1, L0759: 1, L0789: 1, L0759: 1, L0759	AR089: 7, AR316: 6, AR060: 5 L0803: 7, L0766: 4, L0777: 4, L0666: 3, H0521: 3, T0115: 2, H0687: 2, L0799: 2, H0659: 2, L0754: 2, L0779: 2, L0759: 2, S0114: 1, H0341: 1, H0351: 1, S0222: 1, H0441: 1, H0351: 1, H0622: 1, H0909: 1, T0041: 1, H0621: 1, L0371: 1, L0646: 1, L0662: 1, L0774: 1, L0805: 1, L0663: 1, L0659: 1, L0663: 1, L0664: 1, L0651: 1, H0144: 1, T0068: 1, H0519: 1, H0642: 1, L0780: 1, L0748: 1, L0744: 1, L0780: 1, L0758: 1, H0444: 1, L0780: 1, L0758: 1, H0444: 1, H0445: 1, L0590: 1, L0594: 1, S0126: 1 and H0422: 1.	AR089: 194, AR316: 136, AR060: 84 H0251: 7, L0759: 2, H0657: 1, S0116: 1, S0442: 1, S0360: 1, H0486: 1, H0252: 1, S0440: 1, L0764: 1, L0766: 1, L0665: 1, H0144: 1, L0591: 1, L0608: 1 and H0543: 1.	AR089: 40, AR316: 31, AR060: 24
		Lys-22 to His-27, Ser-31 to Gly-38.	
	2927	2928	2929
	1669 - 1884	498 - 632	965 - 609
	330	331	332
	831942	733860	701973
	НСDВ032	HCDBW67	HCDBZ31
	320	321	322

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	2: 4	6	25.		öċ	5: 3		· 4	3:	· 8:		2:				5:	31:	 	. <u>6</u>	<u>;</u> <u>e</u> :	·· · · · ·		11	·:0	 	-:- ::-	33:	:-	.: × ×
03: 1 and	76: 10, H003	7251: 7, HUL7 7598: 6, L043 7036: 5, L051)624: 4, S022	752: 4, L059	H0170: 3, S0358: 3, H0550: H0597: 3, S0346: 3, H0050	051: 3, H025	748: 3, L075	20/25: 3, 30031: 3, 2026s. S0412: 3, H0329: 2, H0374	003: 2, L016	,028: 2, H032	763: 2, L052	546: 2, L053)126: 2, L075	758: 2, LU58	H0008: 2, E0411: 1, S0110: S0400: 1, S0360: 1, S0007:	, S6026: 1, H0175:	3441: 1, H04	1, H0455: 1, H0438:	1, H0486: 1, N0009	H0244: 1, L0021: 1, H0196:	1, H0596: 1, H0327: 1 H0545: 1 H0046:	H0150: 1, N0006: 1, H0565	H0567: 1, H0562: 1, L0471:	1, S0050: 1, H0020:	H0510: 1, S0318:	1, S0214: 1, H0428:	l, H0212: 1, H0163:	H0616: 1, H0551: 1, T0067:	0413: 1, S003 0100: 1, S046
H0251: 2, S0003: 1 and H0547: 1.	S0010: 22, H0144: 12, L0439: 12, L0776: 10, H0032	9, H0038: 8, H0231: /, H0171: 6, H0052: 6, L0598: 6, L0438: 6, H0013: 5, S0036: 5, L0519:	1.0747: 5, H0624: 4, S0222: S0414: 4 H0156: 4 1 0740:			, H0024: 3, S0051: 3, H0252:	, H0412: 3, L0748: 3, L0756:	, E07.59: 5, 50051: 5, E0566. , S0412: 3. H0329: 2, H0374:	, S0049: 2, T0003: 2, L0163:	, H0051: 2, S6028: 2, H0328:	2, L0455: 2, L0763: 2, L0521:	, L0768: 2, L0546: 2, L0532:	, H0519: 2, S0126: 2, L0753:	, L0/31: 2, L0/38: 2, L0389: 110008: 2-1-0411: 1-50110	S0400: 1, 50	, –		, Н0392: 1, Н		, H0244: 1, L0	H0194: H0544	H0150: 1, N(, S0388: 1, HC	, S0336: 1, S0	, L0483: 1, HC	, Н0616: 1, Н	i, H0264: 1, H0413: 1, S0038: 1, S0386: 1, H0100: 1, S0464:
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	Gln-7 to Pro-13, Pro-43 to Gly-50,	Glu-54 to Asp-60, Ala-77 to Ser-94, Ser-134 to I en-139																											
	2930 (J 4 0																											
	32 - 451																												
	333											•			•									-					
	571037													•															
	HCDCB03																												
3	323														·														

D9950083.091201

1, H0538: 1, UNKWN: 1, L0520: 1, L0770: 1, L0364: 1, L0774: 1, L0775: 1, L0540: 1, L0529: 1, S0148: 1, L0352: 1, S0350: 1, S0028: 1, L0786: 1, L0779: 1, H0595: 1, L0592: 1, L0608: 1, L0361: 1, S0106: 1	AR060: 308, AR316: 282, AR089: 255 H0556: 2, H0251: 2, L0766: 2, L0752: 2, H0713: 1, H0255: 1, L0717: 1, H0592: 1, H0688: 1, L0794: 1, L0803: 1, L0776: 1, L0659: 1, H0698: 1, S0330: 1, S0380: 1, H0522: 1, L0743: 1, L0439: 1, L0777: 1, L0485: 1 and H0423: 1.	AR060: 5, AR316: 4, AR089: 3 H0251: 3, H0581: 1 and S0003: 1.	AR214: 30, AR222: 26, AR263: 26, AR264: 24, AR217: 23, AR223: 22, AR169: 22, AR235: 21, AR169: 21, AR165: 21, AR164: 20, AR168: 20, AR166: 19, AR266: 19, AR195: 19, AR291: 18, AR207: 18, AR216: 18, AR207: 18, AR163: 18, AR172: 17, AR224: 16, AR172: 17, AR249: 16, AR197: 16, AR162: 16, AR197: 16, AR209: 16, AR261: 15, AR240: 15, AR261: 15, AR240: 15,
1 L L L L L L L L L L L L L L L L L L L	111112	A A S	44444444444
	2931	2932	2933
	46 - 54	216 - 245	668 - 706
	334	335	336
	813504	847004	841041
	HCDCE51	HCDCI42	HCDDB15
	324	325	326

AR225: 15, AR196: 15, AR289: 15, AR312: 14, AR215: 14, AR212: 14, AR192: 14, AR308: 14, AR257: 14, AR218: 14, AR188: 14, AR219: 14, AR199: 14, AR033: 13, AR181: 13, AR283: 13,	13, AR189: 13, AR288: 12, AR281: 12, AR211: 12, AR247: 12, AR247: 12, AR285: 12, AR285: 11, AR285: 11, AR285: 11, AR175:	11, AR053: 11, AR270: 11, AR182: 10, AR193: 10, AR237: 10, AR252: 10, AR253: 10, AR253: 10, AR253: 10, AR265: 10, AR266:	AR233: 9, AR300: 9, AR174: 9, AR299: 9, AR313: 9, AR267: 9, AR271: 9, AR173: 9, AR296: 9, AR178: 9, AR228: 8, AR061: 8, AR254: 8, AR234: 8,
444444		<u> </u>	. a a a a a a

			1	
			00	
			AR233: 7, AR226: 7,	
			AR204: 7, AR179: 7,	
			AR230: 7, AR060: 7,	
			•	•
	•		AR274: 4	
			H0694: 297, H0656: 17,	• *****
	-		L0803: 8, L0157: 7, L0794: 7,	
			L0747: 6, L0439: 5, L0779: 5,	
		-	L0755: 5, H0622: 4, L0766: 4,	
			L0805: 4, H0547: 4, H0539: 4,	
			L0771: 3, L0809: 3, L0786: 3,	
			L0731: 3, L0588: 3, S0424: 3,	
			H0497: 2, L0623: 2, H0251: 2,	
			H0546: 2, H0545: 2, H0024: 2,	
			H0252: 2, H0617: 2, H0529: 2,	
			L0769; 2, L0533; 2, L0804; 2,	
			L0774; 2, L0775; 2, L0776; 2,	
		· 	S0126: 2, H0521: 2, S0406: 2,	
	•		L0744: 2, L0754: 2, L0756: 2,	
			L0596: 2, L0591: 2, L0599: 2,	
			 H0556: 1, S0218: 1, H0583: 1,	
			 H0657: 1, S0212: 1, H0484: 1,	
			S0420: 1, S0442: 1, S0358: 1,	
			S0376: 1, S0408: 1, H0580: 1,	
			H0208: 1, H0393: 1, H0549: 1,	
			H0643: 1, H0559: 1, H0013: 1,	
		_	H0427: 1, H0575: 1, H0706: 1,	
			S0474: 1, H0052: 1, H0009: 1,	
-			H0178: 1, T0003: 1, S0051: 1,	
			H0375: 1, S6028: 1, H0615: 1,	
			H0031: 1, H0644: 1, H0181: 1,	
			H0032: 1, H0673: 1, S0036: 1,	
	-			
			H0633: 1, S0344: 1, L0640: 1,	
			L0763: 1, L0770: 1, L0761: 1,	
			L0800: 1, L0764: 1, L0521: 1,	
			L0768: 1, L0806: 1, L0655: 1,	

L0782: 1, L0544: 1, L0543: 1, L0787: 1, L0666: 1, L0663: 1, L0665: 1, H0648: 1, H0710: 1, H0696: 1, H0631: 1, S3012: 1, S3014: 1, L0748: 1, L0777: 1, L0758: 1, L0592: 1, L0362: 1, L0601: 1, H0542: 1, H0543: 1 and H0352: 1.	H0251: 6, L0770: 2, L0762: 1 and L0757: 1.	H0136: 2, S0045: 1, H0251: 1, H0413: 1, L0646: 1, L0766: 1 and L0756: 1.	L0748: 11, L0742: 10, H0556: 8, H0251: 6, L0769: 6, L0749: 6, L0587: 6, H0265: 5, H0318: 5, H0040: 5, H0144: 5, H0539: 5, L0750: 5, H0009: 4, H0617: 4, L0758: 4, L0601: 4, T0110: 3, H0038: 3, L0768: 3, L0766: 3, L0809: 3, H0341: 2, S0045: 2, S0046: 2, H0013: 2, H0618: 2, S0182: 2, H0083: 2, H0618: 2, L0772: 2, L0791: 2, L0770: 2, L0372: 2, L0791: 2, L0665: 2, H0576: 2, S0027: 2, S0028: 2, L0731: 2, L0608: 1, H0661: 1, H0125: 1, S0358: 1, S0408: 1, H0208: 1, S0132: 1, H0619: 1, L0717: 1, S0222: 1, H0530: 1, H0052: 1, L0738: 1, H0548: 1, H0486: 1, H0054: 1, H0644: 1, L0456: 1, H0135: 1, H0644: 1, L0456: 1, H0053: 1, H0087: 1, H0413: 1, H0059: 1,
	Lys-23 to Lys-48.		Gin-40 to Cys-48.
	2934	2935	2936
	274 - 417	75 - 170	266 - 412
	337	338	339
	\rightarrow	892137	587264
	HCDDX81	HCDDY28	HCDEB19
	327	328	329

H0494: 1, H0561: 1, L0598: 1, H0529: 1, L0763: 1, L0764: 1, L0773: 1, L0662: 1, L0381: 1, L0774: 1, L0375: 1, L0378: 1, L0805: 1, L0655: 1, L0661: 1, L0527: 1, L0656: 1, L0664: 1, L0382: 1, L0666: 1, L0664: 1, H0593: 1, H0435: 1, H0670: 1, H0660: 1, H0648: 1, H0626: 1, S3014: 1, L0744: 1, L0747: 1, H0595: 1, S0436: 1, L0596: 1, L0605: 1, H0543: 1 and S0424: 1.	AR089: 91, AR316: 66, AR060: 43 L0439: 13, L0747: 13, L0731: 8, L0438: 6, L0754: 6, L0756: 5, L0755: 5, H0624: 4, H0620: 4, L0803: 4, L0809: 4, L0663: 4, S0152: 4, S0354: 3, L0773: 3, L0666: 3, L0750: 3, L0777: 3, L0752: 3, L0759: 3, S0358: 2, H0393: 2, H0369: 2, H0013: 2, H0575: 2, H0251: 2, H0013: 2, H0575: 2, H0251: 2, H0038: 2, L0770: 2, L0774: 2, L0651: 2, L0805: 2, L0774: 2, L0651: 2, L0805: 2, L0749: 2, L0779: 2, L0779: 2, L0749: 2, L0779: 2, L0757: 2, S0026: 2, H0171: 1, H0662: 1, S0420: 1, S0444: 1, S0046: 1, H0619: 1, H0645: 1, H0411: 1, H0550: 1, H0645: 1, H0178: 1, L0471: 1, S0050: 1, T0010: 1, H0687: 1, H0615: 1, H0644: 1, H0674: 1, H0615: 1, H0644: 1, H0674: 1, H0615: 1, H0690: 1, H0599: 1,
	2937
	83 - 262
	340
	838592
	HCDEN46
	330

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	271 - 276	101 - 235	106 - 222
	341	342	343
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	Thr-23 to Lys-28.	Årg-30 to Gly-45, Ser-52 to Ser-65, Gly-101 to Pro-106.
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	23 - 145	374 - 775
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	13
1, S0126: 1, H0684: 1, H0660: 1, H0539: 1, S0152: 1, H0436: 1, L0742: 1, L0740: 1, L0752: 1, L0755: 1, L0757: 1, H0445: 1, L0596: 1, L0592: 1, L0608: 1, S0026: 1, S0194: 1 and H0543: 1.	L0748: 9, L0747: 8, L0759: 7, L0766: 6, L0776: 6, L0665: 5, L0749: 5, H0550: 4, H0052: 4, H0046: 4, L0771: 4, L0741: 2, L0646: 2, L0764: 2, L0741: 2, L0646: 2, S0040: 1, S0358: 1, S0360: 1, H0637: 1, H0637: 1, H0637: 1, H0637: 1, H0637: 1, H0639: 1, H0648: 1, H0639: 1, H0639: 1, H0639: 1, H0659: 1, H0699: 1, H0688: 1, H0428: 1, H0099: 1, H0688: 1, H0428: 1, H0040: 1, H0663: 1, L0764: 1, L0770: 1, L0776: 1, L0776: 1, L0776: 1, L0376: 1, L0376: 1, L0809: 1, L0668: 1, L0778: 1, L0809: 1, H0688: 1, H0653: 1, L0769: 1, L0666: 1, L0778: 1, L0809: 1, H0688: 1, H0659: 1, L0772: 1, H0639: 1, L0772: 1, H0639: 1, L0772:
	7 8 4 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8
	2960
	814 - 996
	363
	874256
	HCE4H32
	353

	•	114290, 138033, 162100, 170500, 170500, 170500, 180860, 264470
		17925
1, S0026: 1, S0194: 1 and H0543: 1.	L0758: 8, L0748: 7, L0764: 5, L0754: 5, L0754: 5, L0775: 3, H0052: 4, H0150: 3, L0777: 3, H0052: 2, L0766: 2, L0771: 2, L0768: 2, L0766: 2, L0774: 2, L0775: 3, H0059: 2, L0776: 2, L0596: 2, L0774: 2, L0775: 1, L0615: 1, H0265: 1, H0657: 1, H0657: 1, H0172: 1, H0687: 1, H0172: 1, H0687: 1, H0040: 1, H0063: 1, L0648: 1, L0648: 1, L0648: 1, L0648: 1, L0657: 1, H0658: 1, H0657: 1, H0658: 1, H0670: 1, H0539: 1, L0658: 1, L0777: 1, L0758: 1, L0777: 1, L0758: 1, L0777: 1, L0758: 1, L0778:	AR089: 15, AR316: 11, AR060: 7 L0601: 17, L0747: 11, H0255: 10, L0766: 10, L0659: 9, H0521: 9, L0439: 9, H0457: 8, L0750: 8, H0543: 8, H0052: 7, L0665: 7, L0731: 7, L0665: 7, L0731: 7, L0666: 5, H0617: 5, H0445: 5, H0656: 5, L0740: 5, H0445: 5, H0656: 4, H0550: 4, H0550: 4, H0550: 4, H0550: 4, H0550: 4, H0650: 4, H0660: 4, H0650: 4, H0650: 4, H0650: 4, H0660: 4, H0650: 4, H0650: 4, H0660: 4, H0660: 4, H0650: 4, H0660: 4, H
1, S0026: H0543: 1	5, L07; 6, H0 1, L06; 1, L0	AR089: AR060: L0601: H0255: 9, H055 7, L066; 7, L066; 5, H061 4, H062 4, H064 4, H064
	His-4 to Thr-9, Ala-31 to Pro-36.	Ser-21 to Trp-32.
	2961	2962
	91 - 321	286 - 384
	364	365
	566797	792953
	HCE4T64	HCE4W88
	354	355

			,
4, L0593: 4, L0595: 4, H0650: 3, H0341: 3, H0484: 3, S0278: 3, H0575: 3, H0251: 3, S0250: 3, H0581: 3, H0221: 3, S0250: 3, T0042: 3, S0210: 3, L0769: 3, L0769: 3, H0542: 3, H0225: 2, H0657: 2, H0483: 2, S0250: 3, H0625: 2, H0657: 2, H0483: 3, H0225: 2, H0657: 2, H06	2, 30539: 2, H0013: 2, L0717: 2, H0261: 2, H0370: 2, H0486: 2, H0205: 2, H0618: 2, H08486: 2, H0009: 2, L0471: 2, H0133: 2, H0264: 2, S0440: 2, H0509: 2, L0767: 2, L0649: 2, L0648: 2, L0378: 2, L0653: 2, L0655: 3, L0656: 3, L0	L0664: 2, L0565: 2, L0689: L0664: 2, L0565: 2, H0689: H0518: 2, S0152: 2, S0206: L0741: 2, L0743: 2, L0748: L0758: 2, L0759: 2, H0707: L0596: 2, L0605: 2, L0361: S0192: 2, H0506: 2, H0265: H0556: 1, H0224: 1, S0110: S0001: 1, H0254: 1, H0663: H0306: 1, H0125: 1, S0356: S0376: 1, S0376: 1	1, H0609: 1, H0586: 1, H0333: 1, H0427: 1, H0590: 1, S0049: 1, H0194: 1, H0309: 1, H0544: 1, H0041: 1, H0024: 1, S0388: 1, H0067: 1, H0271: 1, H0687: 1, H0688: 1, H0428: 1, H0615: 1, H0688: 1, H0181: 1, H0068: 1, H0598: 1, H0181: 1, H06834: 1, H0598: 1, H0181: 1, H0634: 1, H0684: 1, H0684: 1, H0688: 1, H0181: 1, H0688: 1, H0089: 1, H0688: 1, H0087: 1, H0181: 1, H0688: 1, H0087: 1, H0181: 1, H0677: 1, H0181: 1, H0
 <u> 4 でででででででで</u> 刀 班 班 班 班 け 刀 班 班 9 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	v cv	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	3 H H H H H H H H H H H H H H H H H H H

D955023.091501

		147200, 178640, 216900
		2p12
1, H0100: 1, T0041: 1, H0279: 1, H0334: 1, H0625: 1, H0561: 1, S0142: 1, S0426: 1, L0739: 1, S0426: 1, L0639: 1, L0667: 1, L0667: 1, L0673: 1, L0673: 1, L0658: 1, L0638: 1, L0375: 1, L0658: 1, L0376: 1, L0658: 1, L0382: 1, H0648: 1, H0668: 1, H0668: 1, H0668: 1, H0668: 1, H0668: 1, S0424: 1, S0428: 1, and L0600: 1.	H0052: 1	AR089: 17, AR316: 15, AR060: 12 H0617: 7, L0750: 7, H0556: 5, L0769: 3, L0761: 3, L0741: 4, S0132: 3, L0761: 3, L0742: 3, H0618: 2, H0620: 2, H0670: 2, S0328: 2, S0380: 2, L0777: 2, L0771: 2, L0662: 2, L0659: 2, L0771: 2, L0762: 2, H0670: 2, S0328: 2, S0380: 2, L0747: 2, L0753: 2, L0731: 2, H0657: 1, H0656: 1, H0294: 1, H0648: 1, H0656: 1, H0549: 1, H0650: 1, H0649: 1, H0650: 1, H0649: 1, H0650: 1, H0640: 1, H0156: 1, H0640: 1, H0156: 1, H0640: 1, H0156: 1, H0640: 1, H0150: 1, H0640: 1, H0188: 1, H0288: 1, S0250: 1, H0428: 1, H0135: 1, H0135: 1, H0138: 1, H013
	·	Leu-30 to Gly-35. AI AI F 5, 6, 7, 7, 1, 1, 1, 1, 1, 1, 1, 1
	2963	2964
	204 - 293	288 - 482
	366	367
	566864	847032
	HCE5B62	HCE5H86
	356	357

logica" cacused

		309610 309610 309610	192340, 234200
		Xp11	20p13-p12.2
1, H0163: 1, H0090: 1, H0616. 1, T0004: 1, S0438: 1, L0770: 1, L0372: 1, L0646: 1, L0521: 1, L0768: 1, L0766: 1, L5574: 1, L0774: 1, L0775: 1, L0375: 1, L0658: 1, L0540: 1, L0657: 1, L0658: 1, L0540: 1, L0384: 1, L0672: 1, S0188: 1, S0406: 1, H0672: 1, S0188: 1, S0406: 1, H0436: 1, L0779: 1, L0757: 1 and H0506: 1.	H0052: 1 and H0316: 1.	AR060: 4, AR089: 4, AR251: 3, AR249: 3, AR316: 3, AR249: 3, AR274: 2, AR310: 2, AR291: 2, AR269: 2, AR291: 2, AR293: 2, AR175: 2, AR293: 2, AR182: 2, AR293: 2, AR182: 2, AR284: 2, AR313: 1, AR229: 1, AR313: 1, AR312: 1 L0794: 2, L0803: 2, L0754: 2, L0747: 2, H0684: 1, H0085: 1, H0644: 1, H0673: 1, L0760: 1, H0560: 1, L0765: 1, L0774: 1, L0805: 1, L0664: 1, H0689: 1, L0666: 1, L0664: 1, H0689: 1, L0608: 1, L0601: 1, H0542:	L0747: 15, L0748: 11,
		Pro-71 to Leu-83, Pro-119 to Ser-129.	Pro-31 to Pro-37.
	2965	2966	2967
	114 - 233	197 - 586	124 - 324
	368	369	370
	688883	847033	571044
	HCE5164	HCEBF54	HCECO77
	358	329	360

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	118800, 123660, 125660, 125660, 193500, 193500, 193500, 201460, 205100, 237300, 262000, 600266, 601277	
	2435	
L0749: 11, L0731: 10, L0754: 9, L0779: 7, L0777: 7, L0750: 6, H0556: 5, H0547: 5, L0759: 5, H0620: 4, H0135: 4, L0769: 4, L0766: 4, H0135: 4, L0769: 4, L0766: 4, H0135: 4, L0769: 4, L0766: 4, H0660: 4, L0757: 4, L0769: 3, L076: 2, L0774: 2, L0774: 2, L0774: 2, L0774: 2, L0774: 2, L0774: 2, L0776: 1, H0635: 1, H0648: 1, H0638: 1, H0641: 1, H0641: 1, H0672: 1, H0673: 1, L0770: 1, L0662: 1, L0771: 1, L078: 1, L0771: 1, L078: 1, H0642: 1, H0642: 1, H0642: 1, H0642: 1, H065: 1, H0542: 1, H0593: 1, H0593: 1, H0593: 1, H0593: 1, H0506: 1 and H0293: 1, L0780: 1, H0506: 1 and H0293: 1, L0780: 1, H0506: 1 and H0293: 1, L0506: 1 and H0293: 1 a	H0052: 1	L0439: 6, L0777: 5, L0438: 2, L0747: 2, L0756: 2, H0351: 1, H0052: 1, L0663: 1, L0352:
	Lys-36 to Glu-44.	
·	2968	2969
	71 - 238	67 - 207
	371	372
	821321	634525
	нсерн42	HCEDJ05
	361	362

1 and S0412: 1.	L0439: 30, L0740: 12,	L0438: 9, L0742: 9, L0758: 8,	S0022: 6, L0779: 6, S0356: 5,	L0771: 5, L0659: 5, S0360: 4,	L0748: 4, L0592: 4, L0595: 4,	S0242: 4, H0265: 3, S0007: 3,	H0619: 3, S0003: 3, S0214: 3,	H0031: 3, H0488: 3, L0766: 3	L0774: 3, L0805: 3, L0653: 3,	S0126: 3, L0754: 3, L0759: 3,	L0596: 3, L0005: 2, H0097: 2,	H0599: 2, H0318: 2, H0421: 2	H0051: 2, S6028: 2, H0039: 2,	L0142: 2, L0564: 2, L0770: 2,	L0769: 2, L0764: 2, L0649: 2,	L0776: 2, L0666: 2, L0663: 2,	L0665: 2, H0144: 2, H0658: 2,	H0648: 2, S0152: 2, H0522: 2,	L0756: 2, L0777: 2, L0752: 2,	S0260: 2, L0485: 2, L0593: 2,	H0556: 1, S0134: 1, S0116: 1,	S0282: 1, H0662: 1, S0418: 1,	S0420: 1, S0354: 1, S0358: 1,	L0149: 1, S0045: 1, S0140: 1,	H0351: 1, H0550: 1, H0586: 1	H0587: 1, H0574: 1, H0486: 1	H0013: 1, H0244: 1, H0427: 1	H0042: 1, H0575: 1, H0590: 1	S0010: 1, S04/4: 1, S0049: 1,	H0052: 1, H0194: 1, H0309: 1	H059/: 1, H0545: 1, H0046: 1 I 0157: 1 50267: 1 50388: 1	CODS1: 1, 30302: 1, 30368: 1, CODS1: 1 H0365: 1	H0615: 1, H0625: 1, H0269: 1 H0615: 1 H0627: 1 1 0055: 1	H0032: 1, H0169: 1, E0593: 1	S0036: 1, H0591: 1, H0038: 1.	H0616: 1, H0087: 1, H0272: 1	S0038: 1, H0494: 1, H0561: 1,	30210. 1, 110367. 1, 20106. 1,
	2970																																					
	1004 - 1009											-											,,															
	373																																					
	840397												-				•	-															-					
	HCEDJ26																																					
	363																						-															

D99500B3 ..O91201

		176261, 601399		157900, 600631	
		21q22.2		13q13	
L0637: 1, L0772: 1, L0372: 1, L0644: 1, L0521: 1, L0803: 1, L0775: 1, L0376: 1, L0657: 1, L0515: 1, L0526: 1, L0788: 1, H0682: 1, H0543: 1, H0539: 1, H0548: 1, L0751: 1, S0392: 1, L0599: 1, L0594: 1, L0601: 1, L0603: 1, L0366: 1, S0026: 1, S0192: 1, S0196: 1, H0543: 1, H0423: 1 and H0506: 1.	L0748: 3, H0170: 1, H0575: 1, H0052: 1, H0039: 1 and L0793: 1.	AR060: 5, AR316: 5, AR089: 4 L0438: 6, L0439: 6, L0754: 5, H0422: 2, H0638: 1, H0600: 1, H0581: 1, H0434: 1, H0052: 1, S0051: 1, T0010: 1, L0055: 1, H0032: 1, S0386: 1, L0520: 1, L0770: 1, L0521: 1, L0803: 1, L0375: 1, L0518: 1, L0793: 1, L0666: 1, L0665: 1, H0436: 1, H0683: 1, H0522: 1, H0436: 1, L0741: 1, L0751: 1, L0777: 1, L0686: 1 and L0608: 1.	AR089: 32, AR316: 24, AR060: 16 H0052: 2, H0261: 1, H0257: 1, L0657: 1, L0790: 1 and L0777: 1.	L0439: 5, L0779: 2, L0777: 2, L0759: 2, H0156: 1, S0010: 1, S0049: 1, H0052: 1, H0328: 1, L0794: 1, L0804: 1, H0144: 1, L0438: 1, L0352: 1, L0758: 1, L0589: 1 and S0412: 1.	AR060: 4, AR316: 3,
	Ser-15 to Asp-22, Leu-26 to Ala-33.				
	2971	2972	2973	2974	2975
	126 - 248	608 - 712	242 - 307	244 - 351	276 - 281
	374	375	376	377	378
	847034	824312	889968	821322	668234
	HCEDN07	НСЕБО17	HCEEG48	НСЕЕМ33	HCEEP16
	364	365	366	367	368

DOSECOL COLUCTOR

					123100, 153880, 180104, 601649	
					7p21.3-p15.1	
AR089: 2 H0052: 1, H0591: 1, H0132: 1, H0131: 1, S0422: 1, H0659: 1, L0779: 1 and L0780: 1.	H0599: 2, L0439: 2, H0052: 1, L0639: 1, L0792: 1 and S0328: 1.	L0517: 2, S0114: 1, H0656: 1, S0474: 1, H0052: 1, H0179: 1, H0416: 1, H0169: 1, H0538: 1 and 1,0758: 1	H0052: 1	H0052: 1	AR060: 7, AR316: 5, AR089: 2 S0003: 27, L0766: 7, L0438: 5, L0745: 5, S0214: 3, L0439: 3, L0754: 3, L0731: 3, H0635: 2, L0803: 2, L0776: 2, H0144: 2, H0519: 2, L0746: 2, L0756: 2, L0759: 2, L0592: 2, S0400: 1, H0631: 1, H0442: 1, H0574: 1, H0013: 1, H0098: 1, H0056: 1, H0014: 1, H0510: 1, S6028: 1, H0031: 1, H0551: 1, H0541: 1, H0032: 1, S0036: 1, H0591: 1, H0634: 1, L0794: 1, L0774: 1, L0775: 1, L0607: 1, L0659: 1, L075: 1, L0688: 1, H0555: 1, H0518: 1, S0152: 1, H0555: 1, H0518: 1, S0152: 1, H0555: 1, L0752: 1, L0684: 1, S0026: 1, L0752: 1, L0684: 1, S0026:	AR089: 18, AR316: 13, AR060: 8 H0052: 1
					Thr-33 to Trp-38.	
	2976	2977	2978	2979	2980	2981
	18 - 104	282 - 296	223 - 228	232 - 285	224 - 340	166 - 207
:	379	380	381	382	383	384
	826016	996609	695707	822850	637531	745400
	HCEER60	HCEFA10	HCEFA50	HCEFA94	HOEFC27	HCEFG93
	369	370	371	372	373	374

ngsida: dsiel

	203800, 602404
	2p13
H0545: 30, H0673: 13, S0436: 12, H0544: 11, L0731: 8, S0356: 5, H0544: 11, L0731: 8, L0651: 5, L0659: 4, L0659: 4, H0124: 3, L0651: 5, L0771: 3, L0666: 3, L0518: 3, L0771: 3, L0666: 3, L0747: 3, H0717: 2, H0661: 2, S0360: 2, H0550: 2, H00524: 2, L0633: 2, L0803: 2, L0804: 2, L0633: 2, H0696: 2, L0751: 2, H0713: 1, H0662: 1, H04022: 1, H0549: 1, H0369: 1, H0369: 1, H0369: 1, H0369: 1, H0369: 1, H0042: 1, H0039: 1, H0652: 1, H0166: 1, H0169: 1, H0165: 1, H0166: 1, H0169: 1, H0165: 1, H0166: 1, L0644: 1, L0648: 1, L0761: 1, L0647: 1, L0761: 1, L0647: 1, L048: 1, L0375: 1, H0660: 1, S0328: 1, H0662: 1, H0684: 1, L0761: 1,	H0052: 11, H0617: 11, L0741: 9, L0747: 8, H0618: 7, L0439: 7, H0305: 6, L0438: 6, L0659: 5, S0040: 4, S0358: 4, S0222: 4, H0599: 4, H0046: 4, H0050: 4, L010: 4, L0769: 4, L0662: 4, L076: 4, L0774: 4, L0753: 4, L0759: 4, L0005: 3, H0370: 3, H0253: 3, H0181: 3, H0038: 3, H0616: 3, S0038: 3,
Pro-6 to Gln-11, Pro-32 to Arg-37, His-41 to Trp-46, Pro-49 to Gly-54.	Lys-21 to Lys-26, Glu-31 to Gly-38, Leu-184 to Ile-191, Gln-193 to Asp-206, Pro-278 to Leu-286, Gln-305 to Pro-319, Ala-380 to Glu-397, Arg-403 to Thr-412, Leu-423 to Thr-441, Thr-590 to Lys-595,
2982	2983
260 - 439	79 - 2091
382	386
068108	872554
	HCEFK36
375	376

																																_			
75: 3,	77: 3,	58: 3, 62: 2.	19: 2,	36: 2,	98. 2,	38: 2,	2. 5,	48: 2,	88: 2,	42: 2,	02: 1,	12: 1,	56: 1,	30: 1,	17: 1,	86: 1,	.27: 1,	10: 1,	51: 1,	55: 1,	06: 1,	56: 1,	66: 1,	50: 1,	38: 1,	72: 1,	54: 1,	58: 1,	78: 1,	7:1,	17: 1,	55: 1,	26: 1,	59: 1,	36: 1,
3, L07	3, L07	3, L07: 2. H06	2, S004	: 2, S00	. 2, not	2, L06	2, L08(: 2, L07	2, L05	2, H05	: 1, T00	: 1, S02	1, S03	1, H058	1, L07	: 1, H05	: 1, H04	: 1, S00	: 1, H00	1, H035	: 1, T00	: 1, L04	: 1, S03	1, H05	1, H05	1, L077	1, L076	1, L076	1, L0378:	1, L0517:	1, L0647:	1, L066	1, S012	1, H06	1, H04,
L0351: 3, L0794: 3, L0775: 3, L0666: 3, L0663: 3, L0744: 3.	.0740: 3, L0779: 3, L0777: 3,	.0731: 3, L0757: 3, L0758: 3, .0366: 3, H0255: 2, H0662: 2.	S0418: 2, S0420: 2, S0049: 2,	H0327: 2, H0428: 2, S0036: 2,	H0100: 2, H0494: 2, 1,0598: 2,	.0762: 2, L0770: 2, L0638: 2,	J0372: 2, L0767: 2, L0809: 2,	L0664: 2, H0658: 2, L0748: 2,	L0755: 2, L0596: 2, L0588: 2,	H0136: 2, S0276: 2, H0542:	H0265: 1, H0556: 1, T0002:	H0295: 1, H0650: 1, S0212:	S0282: 1, H0589: 1, S0356: 1	S0354: 1, S0360: 1, H0580: 1	S0046: 1, H0393: 1, L0717: 1	H0411: 1, H0438: 1, H0586:	H0587: 1, H0485: 1, H0427: 1	H0156: 1, H0122: 1, S0010: 1,	H0194: 1, H0597: 1, H0051: 1	S0388: 1, S0051: 1, H0355: 1	H0266: 1, H0688: 1, T0006:	H0606: 1, H0674: 1, L0456:	H0124: 1, H0068: 1, S0366:	T0004: 1, L0564: 1, H0560:	L0065: 1, S0150: 1, H0538:	L0637: 1, L0772:	L0800: 1, L0641: 1, L0764: 1,	L0648: 1, L0521: 1, L0768:	L0375: 1,	L0658:	L0519:	L0787: 1, L0790: 1, L0665: 1,	H0520: 1, H0519: 1, S0126: 1	H0690: 1, H0435: 1, H0659: 1	S0044: 1, H0134: 1, H0436: 1, S0037: 1 1 0743: 1 1 0751: 1
351: 3, 3666: 3,	0740: 3,	0731: 3, 0366: 3,)418: 2,	0327: 2	0100:2	762: 2,	3372: 2,	3664: 2,	0755: 2,	0136: 2,	0265: 1,	0295: 1,)282: 1,	354: 1,	046: 1,	0411: 1,	0587: 1,	0156: 1,	0194: 1,	388: 1,	3266: 1,	3606: 1,	0124: 1,	0004: 1,	065: 1,	L0371: 1,	800: 1,	648: 1,	.0803: 1,	.0657: 1,	L0783: 1,	787: 1,)520: 1,)690: 1,	044: 1, 037: 1
77	<u> </u>	<u> </u>	S	Ι,		<u>: </u>	<u> </u>	<u> </u>	<u> </u>	王	Ħ	<u> </u>	<u>x</u>	<u>x</u>	SC	Ħ	田	王	<u> </u>	<u> </u>	Ħ	Ħ	<u>Ħ</u>	<u>T</u>	<u> </u>	<u> </u>	<u> </u>	<u> </u>	<u> </u>	<u> </u>	<u> </u>	<u> </u>	H	H	<u> </u>
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Thr-602 to Gly-608, Ala-626 to Asp-633.	•																																		
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L0754: 1, L0745: 1, L0752: 1, S0031: 1, S0260: 1, S0394: 1, L0584: 1, L0599: 1, L0601: 1, S0192: 1, S0194: 1, H0543: 1, H0423: 1, H0550: 1 and H0352: 1.	L0752: 19, L0439: 12, L0752: 9, L0596: 9, L0759: 6, H0013: 5, H0090: 5, L0662: 5, L0766: 5, H0682: 5, H0436: 5, L0766: 5, H0682: 5, H0436: 5, L0756: 5, L0483: 4, H0553: 4, H0591: 4, L0775: 4, L0362: 4, H06423: 4, H0556: 3, L0776: 3, L0809: 3, H0156: 3, L0777: 3, H0569: 3, H0435: 3, H0659: 3, S00028: 3, L0747: 3, L0777: 3, H0560: 2, S0010: 2, H0318: 2, H0560: 2, S0210: 2, L0761: 2, L0771: 2, L0774: 2, L0805: 2, L0771: 2, L0774: 2, L0805: 2, H0560: 2, S0210: 2, L0761: 2, H0560: 2, S0210: 2, L0761: 2, H0573: 2, L0659: 2, L0788: 2, H0560: 1, H0560: 1, H0571: 1, H0569: 1, H0560: 1, H0571: 1, H0569: 1, H0590: 1, H0635: 1, H0569: 1, H0147: 1, S0346: 1, S0360: 1, H0590: 1, H0564: 1, H0569: 1, H0569: 1, H0564: 1, H0569: 1, H0569: 1, H0564: 1, H0569: 1, H0590: 1, H0575: 1, S0028: 1, H0650: 1, H0564: 1, H0664: 1, H0599: 1, H0564: 1, H0664: 1, H0599: 1,
	Lys-31 to Asn-39.
	2984
	113 - 250
	839253 387
	HCEFN51 8
	377

H0561: 1, S0440: 1, S0150: 1, H0130: 1, L0625: 1, L0770: 1, L0373: 1, L0646: 1, L0764: 1, L0773: 1, L0767: 1, L0364: 1, L0803: 1, L0650: 1, L0375: 1, L0657: 1, L0520: 1, L0529: 1, L0667: 1, L0790: 1, L0793: 1, L0666: 1, L0664: 1, L0665: 1, S0053: 1, H0701: 1, L0352: 1, H0593: 1, H0648: 1, H0671: 1, H058: 1, H0648: 1, H0672: 1, S0330: 1, S0146: 1, H0576: 1, S0330: 1, S0146: 1, H0576: 1, S0390: 1, S0446: 1, L0744: 1, L0746: 1, L0779: 1, L0731: 1, S0260: 1, S0434: 1, S0011: 1,	AR060: 5, AR316: 4, AR089: 4 L0439: 15, H0052: 11, S0007: 9, L0438: 6, L0731: 6, L0779: 5, L0754: 4, H0550: 3, L0769: 3, S0126: 3, L0743: 3, H0194: 2, H0687: 2, H0623: 2, L0768: 2, L0776: 2, L0659: 2, L0666: 2, L0663: 2, H0689: 2, S0330: 2, L0748: 2, L0786: 2, L0777: 2, L0752: 2, L0786: 2, L0777: 2, L0752: 2, L0786: 1, S0336: 1, S0354: 1, S0444: 1, S0356: 1, S0354: 1, S0444: 1, H0431: 1, H0333: 1, H0642: 1, H0575: 1, H0024: 1, S0050: 1, H0150: 1, H0024: 1, L0770: 1, L0372: 1, L0661: 1, L0770: 1, L0794: 1, L0803: 1, L0775: 1, L0653: 1, L0657: 1, L0809: 1,
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JJHSJH	Gly-31 to Gly-46. 3, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,		1,	Cys-3 to His-22. 4, 3, 3, 4, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
	2986	2987	2988	2989
	203 - 373	72 - 254	145 - 330	84 - 179
	389	390	391	392
	847036	844452	846298	658670
	HCEGH74	HCEGK81	HCEGS49	HCEGU75
	379	380	381	382

													•								-							•	
1, H0539: 1, H0522: 1, H0696: 1, S3012: 1, S3014: 1, L0748: 1, L0777: 1, H0543: 1 and H0506: 1	51, AR265: 31, AR309:	28, AR312:	AR053: 26, AR2/3: 21, AR096: 21, AR052: 20,	AR219: 19, AR213: 19, AP247: 18, AP183: 18	18, AR243:	18, AR218:		AR292: 13, AR249: 13, AR282: 13, AR299: 13	12, AR271:	12, AR256:	12, AR283:	11, AR293:	AK186: 10, AK1/9: 10, AR766: 10 AR185: 10	10, AR275:	9, AR300:	9, AR267:	8, AR289:	8, AR296:	AR294: 8, AR231: 8,	o, AK196: 7, AR033:	7, AR280:	7, AR258:	7, AR268:		6, AR226:			AR315: 6, AR270: 6,	0, AR230.
	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	VIV.)
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		Xq28	
AR237: 5, AR061: 5, AR184: 5, AR277: 5, AR206: 5, AR039: 5, AR205: 4, AR194: 4, AR202: 3, AR227: 3, AR204: 3, AR233: 3, AR204: 3, AR281: 2 H0581: 2, L0749: 2, L0779: 2, L0758: 2, H0341: 1, S0376: 1, S0046: 1, H0392: 1, H0590: 1, H0052: 1, L0376: 1, L0515: 1, L0791: 1, H0539: 1 and L0601: 1.	AR089: 8, AR316: 7, AR060: 5 H0052: 1	L0439: 5, L0731: 3, L0769: 2, L0766: 2, L0766: 2, L0649: 2, L0783: 2, L0519: 2, L0751: 2, L0757: 2, L0485: 2, H0265: 1, S0354: 1, S0358: 1, S0376: 1, S0366: 1, H0052: 1, H0046: 1, H0039: 1, S0366: 1, H0135: 1, H0494: 1, H0647: 1, S0344: 1, H0538: 1, L0770: 1, L0772: 1, L0764: 1, L0775: 1, L0776: 1, H0547: 1, L0776: 1, H0547: 1, L0758: 1, L0752: 1, L0758: 1 and H0352: 1.	AR060: 6, AR316: 5, AR089: 4 L0747: 6, L0755: 4, L0759:
		Trp-114 to Thr-131.	Lys-2 to Gln-7.
	2991	2992	2993
	29 - 49	175 - 624	20 - 91
·	394	395	396
	560610	722208	637533
	HCEHW24		НСЕЈР93
	384	385	386

	182600, 186880, 190195, 190195, 190195, 190195, 6022700, 602279, 602279, 602279, 602279, 60279
	14q21.3
4, L0769: 3, L0771: 3, L0439: 3, L0777: 3, H0052: 2, H0266: 2, L0766: 2, L0375: 2, L0740: 2, L0749: 2, L0749: 2, L0749: 2, L0752: 2, L0749: 2, L0749: 2, L0752: 2, L0589: 1, H0458: 1, H0458: 1, H0581: 1, H0581: 1, H0687: 1, H0138: 1, H0135: 1, L0764: 1, L0770: 1, L0643: 1, L0764: 1, L0774: 1, L0763: 1, L0764: 1, L0774: 1, L0763: 1, H0435: 1, L0768: 1, L0768: 1, L0768: 1, L0768: 1, L0768: 1, L0768: 1, H0658: 1, L0768: 1, L07	AR227: 282, AR229: 256, AR226: 235, AR104: 231, AR258: 214, AR061: 209, AR055: 204, AR233: 203, AR237: 200, AR185: 190, AR300: 186, AR186: 183, AR232: 173, AR060: 167, AR234: 164, AR179: 158, AR234: 164, AR179: 125, AR234: 129, AR192: 125, AR299: 124, AR314: 123, AR299: 124, AR314: 123, AR299: 107, AR316: 106, AR299: 107, AR316: 106, AR298: 104, AR231: 102, AR288: 89, AR286: 87, AR206: 88, AR286: 87, AR286: 77, AR248: 87, AR286: 77, AR285: 79, AR280: 77, AR285: 73, AR280: 77, AR285: 73, AR280: 77, AR285: 73,
	Gly-42 to Ala-49.
	108 - 284 2994
	847375 397
	HCELB04
	387

	lpter-p12
AR177: 72, AR291: 71, AR292: 69, AR283: 67, AR295: 67, AR053: 67, AR290: 67, AR218: 65, AR290: 64, AR183: 64, AR290: 64, AR183: 64, AR247: 64, AR089: 62, AR247: 64, AR089: 62, AR247: 59, AR269: 58, AR312: 57, AR281: 56, AR312: 57, AR281: 56, AR213: 55, AR274: 55, AR309: 54, AR251: 54, AR309: 54, AR290: 49, AR277: 48, AR244: 35, AR257: 49, AR310: 44, AR277: 48, AR246: 45, AR257: 42, AR310: 44, AR277: 43, AR290: 28, AR253: 44, AR310: 44, AR277: 42, AR194: 1, AR096: 38, AR244: 35, AR265: 34, AR202: 28, AR265: 34, AR309: 3, L0747: 3, L0741: 8, H0052: 4, L0770: 1, S0282: 1, S0049: 1, H0544: 1, H0178: 1, L0717: 1, H0549: 1, S0036: 1, H0051: 1, L0769: 1, L0769: 1, L0769: 1, L0769: 1, L0769: 1, L0769: 1, L0669: 1, L0792: 1, L0669: 1, L0753: 1, L0669: 1, L0753: 1,	L0366: 1. AR060: 37, AR316: 36, AR089: 34
	Gly-24 to Gly-36, Gly-48 to Arg-60.
	2995
	37 - 219
	398
	846468
	HCEMA08
	388

60358: 4, L0759: 4, L0604: H0083: 3, L0747: 3, H0657: H0341: 2, H0484: 2, S0222: H0052: 2, S0364: 2, H0038: H0413: 2, L0768: 2, L0774: L0775: 2, H0658: 2, L0757: L0758: 2, L0588: 2, L0581: S0026: 2, S0114: 1, S0282: H0638: 1, S0444: 1, S0360: S0410: 1, S0045: 1, H0650: H0559: 1, L0622: 1, H0486: T0114: 1, T0082: 1, H0486: H0173: 1, H0213: 1, L0738: H066: 1, H0616: 1, H0509: H0131: 1, S0422: 1, H0529: L0763: 1, L0766: 1, L0764: L0662: 1, L0766: 1, L0764: L0763: 1, L0741: 1, L0742: L0763: 1, L0769: 1, L0763: L0763: 1, L0766: 1, L0731: S0028: 1, L0769: 1, L0731: S0436: 1, L0608: 1, L0601:	16: 5, 8: 2, S6024: 009: 1, H0051: 52: 1, L0794: 90: 1 and	316: 46, 55: 2, H0580:	10038: 1.	10038: 1. 66: 8, L0438: 31: 3, L0803:
S0358: 4, L0759: 4, L0604: 4, H0083: 3, L0747: 3, H0657: 2, H0341: 2, H0484: 2, S0222: 2, H0342: 2, S0364: 2, H0038: 2, H0052: 2, S0364: 2, L0778: 2, L0775: 2, H0658: 2, L0777: 2, L0775: 2, L0588: 2, L0581: 2, S0026: 2, S0114: 1, S0282: 1, H0638: 1, S0444: 1, S0360: 1, S0410: 1, S0445: 1, R0360: 1, S0410: 1, R0629: 1, H0619: 1, H0550: 1, H0173: 1, H0231: 1, L0738: 1, H0173: 1, H0231: 1, L0768: 1, L0741: 1, S0028: 1, L0741: 1, L0754: 1, L0769: 1, L0768: 1, H0136: 1 and H0352: 1.	AR060: 6, AR316: 5, AR089: 3 L0439: 5, L0438: 2, S6024: 1, H0052: 1, H0009: 1, H0051 1, H0424: 1, S0352: 1, L0794: 1, L0803: 1, L0790: 1 and S0106: 1.	AR089: 58, AR316: 46, AR060: 36 H0052: 3, L0355: 2, H0580: 1 and H0014: 1.	H0052: 1 and H0038:	H0052: 1 and H0038: 1. L0439: 10, L0766: 8, L0438: 3, L0747: 3, L0731: 3, L0803:
			Met-1 to Gly-6.	Met-1 to Gly-6.
	2996	2997	2998	3000
	56 - 217	263 - 370	242 - 358	242 - 358 603 - 686
	399	400	401	402
	563013	740746	564906	850521 564499
	HCENN67	HCENQ22	HCEOF01	HCEOF01 HCEON94
	389	390	391	392 393

	·
0756: 0002: 0775: 0665: 0752:	3, 3, 3, 3, 10123: 10100: 0408: 10100: 10251: 10550: 10652: 10653: 10653: 10653: 10653: 10653: 10663: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667: 10667:
43: 2, L 52: 1, S 68: 1, L 64: 1, L 49: 1, I	Hobbal: 56, Ho521: 47, O769: 20, L0794: 20, L0794: 20, L0747: 3, Ho638: 17, Ho522: 13, O494: 11, Ho620: 7, Ho560: L0788: 17, L0770: 6, Ho519: L0788: 6, L0759: 6, Ho123: Ho087: 5, Ho624: 4, S0408: S0045: 4, Ho052: 4, Ho054: 3, Ho059: 4, Ho059: 3, Ho059: 3, Ho059: 3, Ho059: 2, Lo679: 2, Lo667: 2, Lo667: 2, Lo667: 2, Lo667: 2, Lo607: 3, Ho024: 1, Ho055: 1, Ho053: 1, Lo534: 1, Lo539: 1, S0046: S0013: 1, L0777: 1, Ho055: 1, Lo777: 1, Ho057: 1, Ho0
2, L07 :: 2, H00 :: 1, L07 :: 1, L06 :: 1, L06 543: 1.	H0641: 56, H0521: 47, 10769: 20, L0794: 20, L 18, H0638: 17, H0622: 1 10494: 11, H0620: 7, H 1, L0768: 7, L0768: 7, L0768: 7, L0768: 7, L0778: 6, L0759: 6, F 1, L0758: 6, L0759: 6, F 1, L0757: 5, H0624: 4, S 1, L0757: 5, H0624: 4, S 1, L0757: 5, H0624: 4, S 1, H0545: 4, H0135: 4, F 1, H0545: 4, H0135: 4, F 1, H0545: 4, L0807: 3, L0663: 3, H0435: 3, F 1, H0646: 3, L0807: 3, F 1, H0646: 3, L0807: 3, F 1, H0352: 3, H0435: 3, F 1, H0341: 2, S 1, S 1, S 1, S 1, S 1, F 1, F 1, F 1
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	Ile-23 to Lys-33, Gly-46 to Thr-53.		
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	3017	3018
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1, H0665: 1, S0192: 1, S0242: 1, H0543: 1, H0423: 1 and H0422: 1.	H0046: 9, H0529: 9, L0731: 1 8, L0362: 8, L0659: 6, L0742: 6, S0126: 5, S0328: 5, H0521:	5, L0740: 5, L0777: 5, H0543: 5, S0360: 4, S0003: 4, H0622: 4, S0466: 4, L0662: 4, L0766:	4, L0655: 4, H0696: 4, L0759: 4, S0442: 3, S0444: 3, H0590:	3, H0518: 5, S04/4: 5, S0440: 3, L0776: 3, S0374: 3, S0152: 3, H0624: 3, H0657: 3, S0354:	2, S0358: 2, S0376: 2, H0587: 3 H0524: 3 H0524: 3 H0524: 3 H0626: 3 H0626: 3	H0263: 2, H0150: 2, H0375:	H0615: 2, H0553: 2, H0040: H0634: 2, T0067: 2, S0422:	L0764: 2, L0651: 2, L0653:	2, E0763: 2, E0000: 2, E0003: 2, H0144: 2, L0438: 2, H0660:	S0380: 2, L0754: 2, L0745:	L0601: 2, S0011: 2, S0026:	S0242: 2, H0422: 2, H0170: 1.0615: 1. H0265: 1.	H0716: 1, S0218: 1, L0002:	1, H0650: 1, H0656: 1, S0282: 1, H0661: 1, H0662: 1, S0418:	_	, Н0549: 1, Н0592:		H0003: 1, L0022: 1, H0575:	, H0251: 1, H0596:	1, L0040: 1, H0397: 1, H0123: 1, H0620: 1, H0350: 1, H0024:	1, H0014: 1, H0355: 1, H0266:	1, H0267: 1, H0188: 1, S0022:
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	Pro-6 to Ala-11.	Ala-28 to Trp-33, Ser-85 to Gly-98, Pro-117 to Are-124.
·	3022	3024
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	415	416

COSSOCA OSTECL

6, H0435: 6, H0521: 6, L0751: 5, L0752: 5, H0295: 4, H0617: 4, S0440: 4, L0743: 4, L0779: 4, S0358: 3, H0644: 3, H0313: 3, H0494: 3, L0749: 3, L0749: 3, L0779: 3, L0777: 3, L0596: 3, L0601: 3, H0653: 2, H0264: 2, H0646: 2, S0278: 2, H0646: 2, L0659: 2, L0809: 2, L0665: 2, H0658: 2, L0602: 2, S0152: 2, H0558: 2, L0602: 2, S0152: 2, H0558: 1, H0484: 1, H0664: 1, S0356: 1, H0484: 1, H0664: 1, S0356: 1, H0484: 1, H0686: 1, H0290: 1, H0699: 1, H0790: 1, H0699: 1, H0790: 1, H0059: 1, H0123: 1, H0059: 1, L0194: 1, H0181: 1, H0666: 1, L0666: 1, L0663: 1, L0748: 1, L0643: 1, L0664: 1, L0663: 1, L0748: 1, L0664: 1, L0663: 1, L0748: 1, L0665: 1, L0794: 1, H0659: 1, L0666: 1, L0663: 1, L0789: 1, L0666: 1, L0663: 1, H0672: 1, L0804: 1, L0666: 1, L0663: 1, H0672: 1, H0672: 1, L0663: 1, H0672: 1, H0672: 1, H0672: 1, L0603: 1 and H0445: 1, L0603: 1 and H0445: 1, L0603: 1 and		L0041: 1, H0615: 1, S0152: 1 and H0423: 1.	H0423: 1	L0519: 5, L0748: 5, L0754: 1 4, L0794: 3, L0439: 3, H0031:
Pro-150 to Gly-163, Ser-180 to Asp-186, Trp-201 to Ser-214, Gln-222 to Cys-238, Pro-280 to Ser-288, Pro-312 to Arg-319.	Ser-19 to Pro-37, Pro-54 to Cys-61, Pro-68 to Gly-76.			Thr-4 to Lys-15.
	2615	3026	3027	3028
	437 - 742	78 - 170	44 - 226	167 - 316
	2600	429	430	431
	753260	604583	748224	836110
		HCFLL33	HCFLP48	HCFLQ12
		419	420	421

2, H0673: 2, H0551: 2, L0769: 2, L0805: 2, L0663: 2, L0664: 2, L0779: 2, L0777: 2, L0758: 2, L0591: 2, H0265: 1, H0556: 1, S0046: 1, H0619: 1, L0717: 1, H0370: 1, H0497: 1, H0491: 1, H0545: 1, H0650: 1, H0373: 1, T0010: 1, S6028: 1, S0250: 1, H0625: 1, S0440: 1, S0210: 1, L0761: 1, L0771: 1, L0662: 1, L0766: 1, L0579: 1, L0540: 1, L0547: 1, L0518: 1, L0530: 1, L0788: 1, L0790: 1, S0330: 1, S0378: 1, L0601: 1, S0192: 1, H0642: 1 and H0423: 1.	AR089: 8, AR316: 6, AR060: 5 L0766: 6, H0616: 3, L0809: 3, L0789: 3, H0556: 2, S0144: 2, L0770: 2, L0761: 2, L0803: 2, L0757: 2, S0342: 1, S0116: 1, H0341: 1, S0358: 1, H0351: 1, S0278: 1, H0156: 1, H0004: 1, H0251: 1, H0014: 1, T0006: 1, H0424: 1, H0641: 1, L0374: 1, L0771: 1, L0794: 1, L0804: 1, L0774: 1, L0794: 1, L0804: 1, L0774: 1, L075: 1, L0764: 1, H0518: 1, H0522: 1, S0044: 1, H0518: 1, H0522: 1, L0780: 1, H0445: 1, L0750: 1, L0780: 1, H0445: 1, H0543: 1, H0423: 1 and H0677: 1.	316: 6, 16: 3, L0809: 556: 2, S0144:
2, H0673: 2, H0551: 2, 2, L08055: 2, L08055: 2, L0663: 2, 2, L0777: 2, 2, L0777: 2, 2, L0591: 2, H0655: 1, 1, S0046: 1, H06497: 1, 1, H0545: 1, H0640: 1, H0625: 1, S0440: 1, 1, L0761: 1, L0762: 1, and H0423	AR089: 8, AR316: 6, AR060: 5 L0766: 6, H0616: 3, L0809: 3, L0789: 3, H0556: 2, S0144: 2, L0770: 2, L0761: 2, L0803: 2, L0757: 2, S0342: 1, S0116: 1, H0341: 1, S0358: 1, H0351: 1, S0278: 1, H0146: 1, H0004: 1, H0251: 1, H0014: 1, L0374: 1, L0771: 1, L0794: 1, L0804: 1, L0774: 1, L0775: 1, L0376: 1, L0655: 1, L0783: 1, H0683: 1, H0518: 1, H0522: 1, S0044: 1, H0436: 1, L0742: 1, L0754: 1, L0749: 1, L0750: 1, L0754: 1, H0445: 1, H0543: 1, H0423: 1, H0445: 1, H0543: 1, H0423: 1, H0445: 1, H0543: 1, H0423: 1, H0677: 1.	AR089: 8, AR316: 6, AR060: 5 L0766: 6, H0616: 3, L0809: 3, L0789: 3, H0556: 2, S0144:
		Pro-47 to Pro-59, Gly-96 to Asp-101, Arg-114 to Trp-121, Pro-132 to Trp-138.
	3029	3030
	385 - 411	95 - 568
	432	433
	786452	858875
	HCFLY20	HCFLY20
	422	423

2, L0770: 2, L0761: 2, L0803: 2, L0757: 2, S0342: 1, S0116: 1, H0341: 1, S0358: 1, H0351: 1, S0278: 1, H0156: 1, H0004: 1, H0251: 1, H0014: 1, T0006: 1, H0424: 1, H0641: 1, L0374: 1, L0771: 1, L0794: 1, L0804: 1, L0774: 1, L0775: 1, L0376: 1, L0774: 1, L0775: 1, L0376: 1, H0555: 1, L0783: 1, H0683: 1, H0548: 1, L0742: 1, L0754: 1, H0445: 1, H0543: 1, H0423: 1, H0445: 1, H0543: 1, H0423:	: 1	5, AR184: 5, 4, AR269: 4, 4, AR270: 4, 3, AR299: 3, 3, AR299: 3, 3, AR299: 3, 3, AR299: 3, 3, AR299: 2, 2, AR296: 2, 2, AR296: 2, 2, AR296: 2, 2, AR296: 2, 2, AR286: 2, 2, AR288: 2, 2, AR288: 2, 2, AR286: 2, 2, AR300: 2, 2, AR314: 2, 2, AR3232: 2, 2, AR286: 2, 2, AR323: 2, 2, AR323: 2, 2, AR337: 1
2, L0770: 2, L0 2, L0757: 2, S0 1, H0341: 1, S0 1, S0278: 1, H0 1, H0251: 1, H7 1, H0424: 1, H7 1, L0771: 1, L0 1, L0774: 1, L0 1, L0655: 1, L0 1, H0436: 1, L0 1, L0749: 1, L0 1, L0749: 1, L0 1, L0749: 1, L0 1, L0749: 1, L0		
	5 3031 Ser-21 to Ser-28	3032
	434 217	435 321
	\vdash	HCFMJ40 604589
	424	425

		203310					
		6q12-q13					
AR275: 1, AR284: 1, AR263: 1, AR247: 1, AR179: 1, AR039: 1, AR281: 1, AR244: 1, AR259: 1 H0423: 1 H0412: 2, H0187: 2, L0777:	2, S0114: 1, H0306: 1, H0402: 1, S0354: 1, H0333: 1, H0052: 1, H0413: 1, L0770: 1, L0776: 1, H0627: 1, L0750: 1, L0779: 1, L0731: 1 and H0423: 1.	L0766: 2, S0354: 1, S0376: 1, H0574: 1, H0486: 1, L0369: 1, L0637: 1, L0775: 1, L0666: 1, L0665: 1, S0374: 1, L0352: 1, S0330: 1, L0439: 1, L0779: 1, L0758: 1, H0423: 1, S0424: 1 and H0506: 1.	H0423: 1	S0001: 1, S0476: 1, H0264: 1, H0521: 1, H0423: 1 and H0422: 1.	L0758: 7, H0040: 5, L0731: 3, S0444: 2, L0794: 2, H0701: 2, H0583: 1, S0418: 1, S0358: 1, H0486: 1, H0354: 1, H0591: 1, L0803: 1, L0774: 1, L0805: 1, L0665: 1, L0438: 1, L0754: 1 and H0423: 1.	H0306: 1 and H0423: 1.	AR176: 4, AR282: 4, AR053: 4, AR183: 3, AR263: 3, AR250: 3, AR225: 3, AR272: 3, AR197: 3, AR291: 2, AR299: 2, AR182: 2, AR2905: 2, AR182: 2,
3033		3034	3035	3036	3037	3038	3039
110 - 394		112 - 210	354 - 440	191 - 340	448 - 543		183 - 323
436		437	438	439	440	441	442
580817		894836	581042	825989	746864	732010	740934
HCFML07		HCFMR75	HCFMX16	HCFMX88	HCFNM40	Н	HCFNN16
426		427	428	429	430	431	432

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2, AR039	2. AR196:	2. AR214:	AR171	AR195	2, AR199	AR228	, AR261	AR096.	AR283:	AD202	AD170		AR257:	, AR269:	. AR211:	L0740: 22, L0750: 21	7752.	3 6	IS, LU/69: 9, LU/57: 9,	341: ,	083: :	114: 4	687.	7 :689	H0224: 3, H0586: 3, H0135:	771.	1,0776; 3,1,0657; 3, H0659;	521.	171:	420: 7	550: 2	351: 2	038:	625:	763: 2	555: 2	563: 2	751: 2	599: 2	H0423: 2, H0225: 1, H0686:
	2.	, 5, 4	, C	; c;	, 2, A	2, 4	1, 4	\ -	, , , ,	, -	- -	- ,	ή,	_	\neg	22.1	14.	1 6	., y	9, HO	6, HO	4. SO	4 H0	t. H0	3. HO	3.1.0	3,10	3 HO	3. HO	2, S0,	2. HO	2, SO	2, H0	2, H0	2. LO	2, LO	2. LO	2, 10	2, L0	2, H0
AR287:	AR216:	AR060:	AR313.	AR178:	AR308:	AR207:	AR089:	AR312.	AR213:	A D 100.	AD280.	707	AR185:	AR201:	AR297:	0740	747.		מן	758: (775: (170:	157.	126: 4	224:	494:	.911	.859	749.		345: 2	024:	617:	413:	762: 2	554:	783: 2	742: 2	591: 2	423:
AR	AR	AR	AR	A. A.	AR	AR	A.	AR	AR.	<u> </u>	2 2	£ :	AK	<u>A</u>	AR	_		3 4	15,	<u> </u>	<u>C</u>	<u>H</u>	1.0	8	H	9	0,1	Î	0	H	SO.	9	<u>H</u>	H	0	Š		20	<u>~</u>	<u>H</u>
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H0656: 1, S0116: 1, H0671: 1, H0177: 1, H0638: 1, H0125: 1, S0356: 1, H0580: 1, S0046: 1, H0351: 1, H0638: 1, H0125: 1, H0632: 1, L0622: 1, H0648: 1, H0632: 1, L0622: 1, H0486: 1, H0036: 1, H0390: 1, H0196: 1, H0036: 1, H0290: 1, H0196: 1, H0099: 1, H0290: 1, H0290: 1, H0290: 1, H0051: 1, H0619:	AR089: 1 H0580: 1, H0431: 1, S0010: 1, H0379: 1, H0520: 1, H0547: 1, H0435: 1, L0602: 1, L0740:
	AF H H 1, 1,
	3040
	241 - 390
	443
	762959
	HCFNN75
	433

		•																																		
H0423: 1	H0423: 2, L0771: 1, L0768: 1, L075: 1, L0809: 1, H0435:	1, L0779: 1 and H0422: 1.	H0458: 1	AR183: 6, AR055: 5, AR244: 5 AR247: 5) 4	AR089: 4, AR234: 4,	AR309: 3, AR282: 3,		AR231: 3, AR052: 3,		AR053: 3, AR238: 3,	AR267: 3, AR243: 3,	AR269: 3, AR310: 2,			AR237: 2, AR291: 2,		AR295: 2, AR316: 2,	2, AR293:	2, AR292:		2, AR240:	2, AR312:	2, AR286:			(1)	_	AR060: 1, AR300: 1,	AR033: 1, AR289: 1,	AR185: 1, AR218: 1,	AR177: 1, AR205: 1,	AR283: 1, AR194: 1,		AR281: 1, AR039: 1	L0665: 6, L0747: 5, L0743:
				Ala-10 to Arg-18, Glu-45 to Glu-55		,	7	 	7	7	7	7		/	7	7	/	7	7	4	7	4	4	4	7	₫_	\forall	₹	V	A	A	▼	<u> </u>	4	A	
3041	3042		3043	3044																																
27 - 131	3 - 98		83 - 181	201 - 851																					•											
444	445		446	447									-																							
747700	668236		604603	610370				 														••												-		
HCFOG17	нсғон93		\rightarrow	HCHAC68				-						•													•					•				
434	435		436	437																												-				

3, S0420: 2, H0266: 2, H0494: 2, L0646: 2, L0662: 2, L0806: 2, L0659: 2, L0662: 2, L0806: 2, H0295: 1, H0294: 1, H0483: 1, S0132: 1, S0222: 1, H0370: 1, H0600: 1, H0586: 1, H0587: 1, L0021: 1, H0688: 1, H0039: 1, H0622: 1, H0551: 1, H0116: 1, L0763: 1, L0770: 1, L0764: 1, L0771: 1, L0773: 1, L0386: 1, L0803: 1, L0775: 1, L0653: 1, L0663: 1, H0555: 1, L0744: 1, L0779: 1, L0731: 1, L0757: 1, L0605: 1 and H0543: 1.	L0749: 5, \$0126: 4, L0774: 3, L0659: 3, L0789: 3, L0750: 3, \$0420: 2, H0620: 2, L0763: 2, L0764: 2, L0775: 2, \$0404: 2, L0743: 2, L0748: 2, H0624: 1, H0295: 1, \$0134: 1, \$0282: 1, H0483: 1, H0254: 1, H0638: 1, \$0442: 1, \$0376: 1, \$0444: 1, H0675: 1, H0580: 1, H0156: 1, \$0049: 1, H0309: 1, H0009: 1, H0659: 1, H0625: 1, H0625: 1, H0659: 1, H0625: 1, L0643: 1, L0776: 1, L0663: 1, L0665: 1, \$0374: 1, H0435: 1, L0659: 1, H0648: 1, L0439: 1, L0759: 1, H0648: 1, L0439: 1, L0759: 1, L0591: 1 and H0506: 1.	L0769: 4, L0794: 3, L0751: 3, H0659: 2, H0483: 1, S0007: 1, H0018: 1, H0615: 1, L0763: 1, L0638: 1, L0800: 1, L0768: 1, L0522: 1, L0803: 1, L0805: 1, L0776: 1, L0790: 1, L0777: 1 and L0758: 1.
3, S0420: 2 2, L0646: 2 2, L0659: 2 1, R0132: 1 1, H0600: 1, L0021: 1 1, L071: 1 1, L0803: 1 1, L0663: 1, L0770: 1 1, L0663: 1, L0663: 1, L0663: 1	1, 1074 3, 106 3, 106 3, 104 2, 107 1, 108 1, 108 1, 108 1, 108 1, 108 1, 108 1, 108 1, 108 1, 108	1, H00 1, H00 1, L06 1, L05 1, L07 1, L07 1, L07
	3045	3046
	234 - 344	267 - 446
	448	449
	892141	840333
	НСНВР49	HCHCA79
	438	439

H0013: 2, H0036: 2, L0776: 2, H0483: 1, S0045: 1, L0162: 1, H0252: 1, H0048: 1, S0015: 1, S0382: 1, L0766: 1, H0144: 1, S0013: 1, L0740: 1, L0745: 1, L0777: 1, L0759: 1 and L0366: 1.	29, AR202: 24, 23, AR246: 19, AR246: 19, AR266: 19, AR248: 18, AR244: 18, 17, AR289: 17, AR289: 17, AR288: 15, AR206: 13, AR295: 13, AR295: 13, AR295: 13, AR295: 11, AR300: 12, AR300: 11, AR299: 11, AR290: 11, AR290: 11, AR290: 11, AR290: 10,
H0013: 2, H0036: 2, L07 2, H0483: 1, S0045: 1, L0 1, H0252: 1, H0048: 1, S0 1, S0382: 1, L0766: 1, H0 1, S0013: 1, L0740: 1, L07 1, L0777: 1, L0759: 1 and L0366: 1.	AR281: 29, AR202: AR205: 23, AR194: AR039: 20, AR246: AR315: 19, AR263: AR206: 18, AR244: AR243: 17, AR282: AR266: 17, AR282: AR283: 17, AR183: AR241: 15, AR204: AR271: 15, AR204: AR214: 13, AR266: AR271: 13, AR204: AR214: 13, AR206: AR215: 12, AR310: AR215: 12, AR310: AR213: 12, AR300: AR206: 12, AR211: AR213: 11, AR295: AR238: 11, AR299: AR238: 11, AR299: AR238: 11, AR291: AR213: 10, AR293: AR312: 11, AR291: AR312: 11, AR291: AR313: 10, AR293: AR313: 10, AR293: AR309: 9, AR293: AR309: 9, AR293: AR294: 9, AR275: AR294: 9, AR273:
Glu-28 to Lys-34.	Lys-111 to Thr-117, Asn-193 to Trp-199, Ala-271 to Pro-282, Leu-303 to Ala-327.
3047	3048
576 - 743	168 - 1211
450	451
862534	833049
нснсд33	HCHMY57
440	144

			180297, 230450, 248611, 251000, 263200, 600211, 600701, 601690
·			6p21-p12
AR229: 8, AR218: 8, AR293: 8, AR175: 8, AR231: 7, AR256: 7, AR226: 7, AR219: 7, AR258: 7, AR061: 7, AR233: 6, AR060: 6, AR253: 6, AR237: 6, AR186: 6, AR259: 6, AR186: 6, AR259: 6, AR186: 6, AR197: 4, AR248: 5, AR179: 4, AR249: 4 H0484: 3, L0439: 3, S0354: 2, H0521: 2, H0265: 1, H0341: 1, S0356: 1, H0427: 1, H0052: 1, H0494: 1, S0422: 1, L0666: 1, L0438: 1, S0208: 1, L0743: 1, L0749: 1, L0777: 1, H0445: 1, L0749: 1, L0777: 1, H0445: 1, S0436: 1 and H0543: 1.	AR089: 18, AR316: 14, AR060: 10 H0484: 1	L0766: 4, L0591: 4, H0657: 2, H0593: 2, L0759: 2, L0596: 2, H0656: 1, H0484: 1, S0418: 1, S0420: 1, S0356: 1, H0587: 1, H0544: 1, H0135: 1, L0667: 1, L0800: 1, L0771: 1, L0768: 1, L0804: 1, L0803: 1, L0804: 1, L0749: 1, L0749: 1, L0749: 1, L0779: 1 and L0731: 1.	L0748: 9, L0665: 4, H0486: 3, L0761: 3, L0663: 3, L0752: 3, S0116: 2, S0358: 2, L0662: 2, L0659: 2, L0657: 1, H0484: 1, H0402: 1, S0420: 1, H0580:
	7	Ser-4 to Gly-10, Ala-13 to Gly-31, Gln-41 to Lys-68, Gln-73 to Asp-78.	Gly-16 to His-22, Ser-51 to Glu-61. 2 2 2 2
	3049	3050	3051
	341 - 427	24 - 1253	1321 - 1521
	452	453	454
	688042	837297	793648
	900ОНОН	нсноу52	нснов93
	442	443	444

	180297, 230450, 248611, 251000, 263200, 600211, 601690	143890, 151440, 600173, 600276, 600310, 600310, 601843
	6p21-p12	19p13.1
1, H0485: 1, L0738: 1, H0046: 1, H0239: 1, S0022: 1, H0553: 1, H0038: 1, H0551: 1, H0494: 1, H0560: 1, H0509: 1, S0002: 1, L0645: 1, L079: 1, L0766: 1, L0803: 1, L0652: 1, L0657: 1, L0793: 1, H0144: 1, H0547: 1, H0519: 1, H0672: 1, H0539: 1, H0710: 1, H0521: 1, H0696: 1, H0478: 1, L0439: 1, L0777: 1, L0750: 1, L0756: 1, L0757: 1, H0595: 1, H0542: 1, H0422: 1 and L0462: 1.	L0748: 9, L0665: 4, H0486: 3, L0761: 3, L063: 3, L0752: 3, S0116: 2, S0358: 2, L0662: 2, L0659: 1, H0484: 1, H0485: 1, L0738: 1, H0494: 1, H0560: 1, H0551: 1, H0569: 1, L0667: 1, L0645: 1, L0671: 1, L0667: 1, L0645: 1, L0672: 1, H0549: 1, H0519: 1, H0551:	AR089: 46, AR316: 33, AR060: 21 L0740: 15, L0752: 11, L0731: 11, H0052: 9, L0769: 9, L0748: 9, L0753: 8, L0746: 7, L0747: 7, L0775: 6, L0751:
	Gly-16 to His-22, Ser-51 to Glu-61.	
	3052	3053
	1321 - 1521	1050 - 1139
	455	456
	875853	845659
	нснов93	нсг вк 61
	445	446

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	3072	3073	3074	3075
	516 - 593	72 - 191	292 - 321	142 - 276
	475	476	477	478
	862285	832140	845771	834331
	HCQCV23	нсоррэз	нсорр61	нсорт67
	465	466	467	468

DGSSOORS OGLEGI

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			Ala-46 to Arg-62, Leu-68 to Thr-94.
	3076	3077	3078
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	479	480	481
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	HCRAI29	HCRB179	HCRBL20
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		Arg-7 to Ser-28 t Arg-50 1
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Glu-76 to Pro-81, Cys-115 to Val-127, Thr-140 to Asp-171, Ala-175 to Gly-187.	Arg-7 to Pro-13.		
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	41 - 139	86 - 154	2164 - 2340
	2601	483	484
	722210	897005	849071
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	3p22
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H0520: 3, H0521: 3, L0744: 3,	L0745: 3, L0752: 3, H0422: 3, S0424: 3, H0624: 2, H0638: 2	S0424; 3, D0624; 2, D0636; 2, S0360; 2, H0549; 2, S0222; 2,	S0049: 2, H0014: 2, H0051: 2,	S0388: 2, H0266: 2, H0040: 2,	S0210: 2, L0761: 2,	L0667: 2, L0662: 2, L0518: 2,	.0666: 2, L0664: 2, L0665: 2,	S0378: 2, S0152: 2,	H0522: 2, L0356: 2, H0631: 2,	L0743: 2, L0751: 2, L0754: 2,	L0747: 2, L0755: 2, H0445: 2,	L0596; 2, L0588; 2, L0591; 2,	L0608: 2, L0604: 2, L0366: 2,	H0542: 2, H0543: 2, H0423: 2,	H0171: 1, L0442: 1, S0114: 1,	S0212: 1, H0661: 1, L0005: 1,	S0358: 1, H0675: 1, S0132: 1,	S0300: 1, S0278: 1, H0369: 1,	H0441: 1, H0431: 1, H0592: 1,	H0331: 1, S0414: 1, H0486: 1,	S0665: 1, S0346: 1,	H0318: 1, S0474: 1, H0581: 1,	H0421: 1, H0009: 1, N0006: 1,	H0024: 1, S0051: 1, H0271: 1,	H0188: 1, S0318: 1, H0687: 1,	S0214: 1, H0252: 1,	L0194: 1, H0553: 1,	H0628: 1, H0032: 1, H0212: 1,	.0456: 1, H0090: 1, H0616: 1,	H0551: 1, T0067: 1, H0264: 1,	•	_	<u> </u>	_		0766: 1, L0389: 1,	L0388: 1, L0650: 1, L0805: 1, 1, 0683: 1, 1, 0776: 1, 1, 0606: 1	0//U. 1, LOUVU. 1,1
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